Access DB# 6548

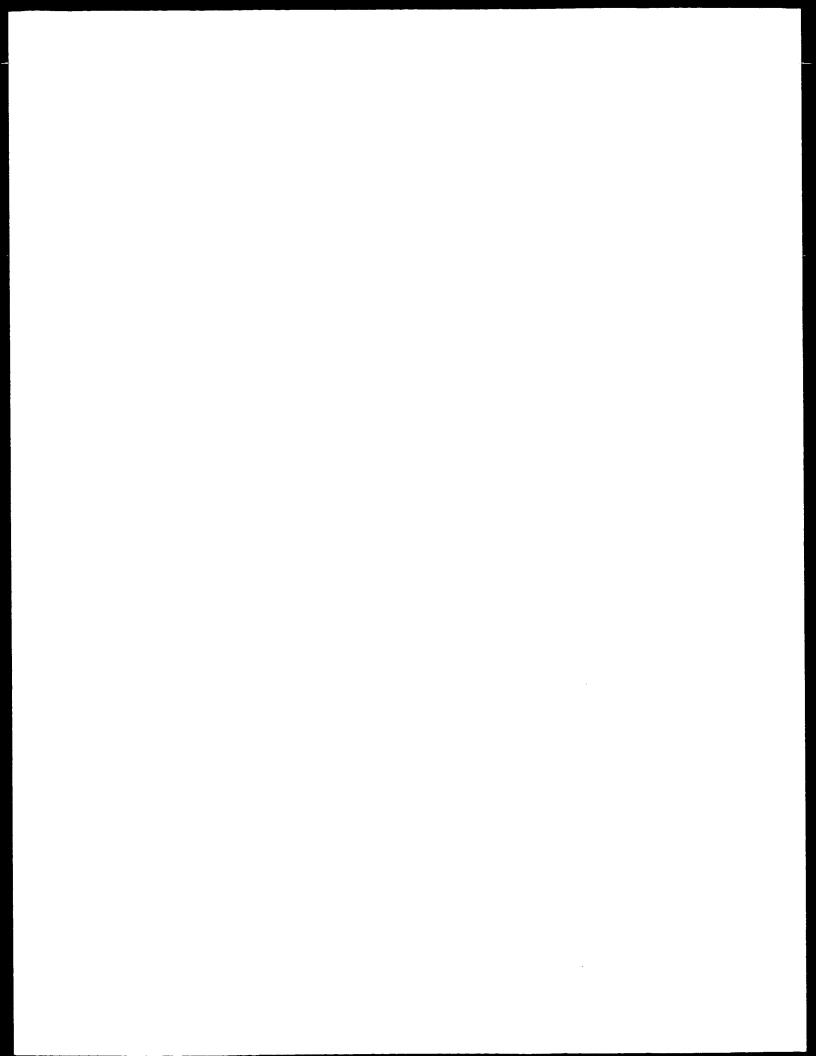
#### SEARCH REQUEST FORM

#### Scientific and Technical Information Center

	scientific and recum	cal information Center
Requester's Full Name: Mall-Art Unit: 1942 Phone Mail Box and Bldg/Room Location  SE12 SA	on: Ke	Examiner #: 15472 Date: 2/12/2003  Serial Number: 05/935440  sults Format Preferred (circle): PAPER DISK E-MAIL
If more than one search is sub-	mitted, please priorit	tize searches in order of need.
Please provide a detailed statement of the Include the elected species or structures,	e search topic, and describ keywords, synonyms, acro is that may have a special r	ne as specifically as possible the subject matter to be searched.  conyms, and registry numbers, and combine with the concept or  meaning. Give examples or relevant citations, authors, etc, if
Title of Invention:		
Inventors (please provide full names):		
(P P III IIII IIII)		
Earliest Priority Filing Date:		
		(parent, child, divisional, or issued patent numbers) along with the
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		historical Librarian historical State Chemical Library CM 1:E07 - 703-308-4493
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STAFF USE ONLY	Type of Search	Vendors and cost where applicable
Searcher:	NA Sequence (#)	STN
Searcher Phone #:	AA Sequence (#)	Dialog
Searcher Location:	Structure (#)	Questel/Orbit
Date Scarcher Picked Up: 2 1 1 1 0 2	Bibliographic	Dr.Link
Date Completed: 31343	Litigation	Lexis/Nexis
Searcher Prep & Review Time:	Fulltext	Sequence Systems
Clerical Prep Time:	Patént Family	WWW/Internet

PTO-1590 (8-01)

Clerical Prep Time:
Online Time:



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Sequence 17, Appl
                                                                                                             Pebruary 12, 2003, 11:12:42 ; Search time 17.4167 Seconds
(without alignments)
32.098 Million cell updates/sec
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Sequence 6,
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3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-081-975-10

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Maximum Match 100%
Listing first 45 summaries
                                                                       OM protein - protein search, using sw model
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1 PPLSQETFSDLWKLLPENG 19
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Gapop 10.0 , Gapext 0.5
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Match Length
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Maximum DB seq length: 25
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Perfect score:
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                                                                                                                    Run on:
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODIES OF TREATMENT USING
TITLE OF INVENTION: ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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COMPUTER: MA
COMPUTER: Diskette
MEDIUM TYPE: Diskette
COMPUTER: ENA Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSED for Windows
SOFTWARE: FastSED for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
FLING DATE: 12-MAY-1998
FLING DATE: 12-MAY-1998
FLING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELEDOMUNICATION INFORMATION:
TELEBHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.3%; Score 99; DB 4; Le
100.0%; Pred. No. 1.6e-09;
tive 0; Mismatches 0;
          US-08-424-957-45

US-09-424-957-46

US-09-035-686-46

US-09-035-686-46

US-09-400-653A-7

US-09-240-061B-7

US-08-277-660A-10

US-08-424-957-28

US-08-424-957-28

US-08-424-957-28

US-09-035-686-28

US-09-035-686-11

US-08-424-957-11

US-09-035-686-11
                                                                                                                                                                                                                                                                                        ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                            Sequence 8, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
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LENGTH: 20 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 94.3
Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 617-345-1300
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TOPOLOGY: linear
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RESULT 4
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APPLICANT: Kaelin, William
APPLICANT: JOSE, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                   APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: Sast, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: BOSTON
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94.3%; Score 99; DB 4; Length 20;

Best Local Similarity 100.0%; Pred. No. 1.6e-09;

Matches 18; Conservative 0; Mismatches 0; Indels
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OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11, Application US/09081975 Patent No. 6451979
                                                                                                                                                     Sequence 10, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 4740
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nixon Peabody LLP
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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CORRESPONDENCE ADDRESS:
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                                                                                                                                    US-09-081-975-10
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US-09-081-975-16
Sequence 16, Application US/09081975
Sequence 16, Application US/09081975
Sequence 16, Application US/09081975
Sequence 16, Application US/09081975
Sequence 16, Application of Association Sequence 16, Application Sequence 17 Title OF INVENTION: METHODS OF TREATMENT USING TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE TITLE OF INVENTION: ANTIBODIES AND PROTEINS THERETO, AND USES OF THE CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.3%; Score 99; DB 4; Length 20; 100.0%; Pred. No. 1.6e-09; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Comparatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
                                                                                                                                                     IBM Compatible
SYSTEM: Windows
FastSEQ for Windows Version 2.0b
                                                                                                                                                                                           CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
TLING DATE: 12-MAY-1998
CLASSIFICATION T435
PRIOR APPLICATION T435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Romald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECHONE: 617-345-6054
TELECHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          101 Federal Street
101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.0
                                                                                                                                    Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear
US-09-081-975-11
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OPERATING SYSTEM:
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STREET: Lor
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                          Boston
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ZIP: 02110
                                                                                       02110
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                                                                  COUNTRY:
    STREET:
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APPLICANT: Kaelin, William
APPLICANT: JOST, Christine
APPLICANT: ADST, Christine
TITLE OF INVENTION: METHOS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
TOTLE OF INVENTION: ANTIBODIES
CORRESPONDENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/09081975

Patent No. 6451979

GENERAL INFORMATION:

APPLICANT: Jost, Christine

TITLE OF INVENTION: METHODS OF TREATMENT USING

TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE

TITLE OF INVENTION: ANTIBODIES

NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SURFACE TEASTSED for Windows Version 2.0b CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                        Sequence 15, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                     1 PPLSQETFSDLWKLLPEN 18
                                                                 2 PPLSQETFSDLWNLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 PPLSQETFSDLWNLLPEN 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 94.4
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: linear
US-09-081-975-15
                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 101 r.
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                   US-09-081-975-15
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US-09-081-975-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE:
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Patent No. 6451979
GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Greet, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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0
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

89.5%; Score 94; DB 4; Length 20;
Best Local Similarity 94.4%; Pred. No. 1e-08;
Matches 17; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compacible
COMPUTER: IBM Compacible
OPERATING SYSTEM: Windows
SOFTWARE: FastEND for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY-AGENT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30,628
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEPRAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                   INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 amino acids
                                                                                                                                                                                                                         ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-081-975-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: ADSC, Christine
TITLE OF INVENTION: METHOD SOF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LiP
STREET: 101 Federal Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPREATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION: 435
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                       MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 415
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19, Application US/09081975 Patent No. 6451979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47400
                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFREENCE/DOCKET NUMBER: 4740(
TELECOMMUNICATION INFORMATION:
TELERPHONE: 617-345-6054
TELEFAX: 617-345-1300
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSOETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-09-081-975-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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STREET: 102
TTV: Boston
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                                                                                         USA
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                                                                                           COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Discussion of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
CONTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                                                                   81.9%; Score 86; DB 4; Length 20; 94.1%; Pred. No. 1.9e-07; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUBER: US/08/277,660A FILING DATE: 20-JUL-1994 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 73.3%; Score 77; DB 1; Le Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A-60244/WHD
                   NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence 1, Application US/08277660A ; Patent No. 5702908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY AGENT ASTER H.
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-6
TELECOMMUNICATION INFORMATION:
TELEFHONE: (415) 781-1989
                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LECORDOR: (415) / C. TELEPHONE: (415) 398-3249
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3 PLSQETFSDLWKLLPPN 19
                                                                                                                                                                                                                                                                                                                                                                                                                      16; Conservative
                                                                                                                                                                                                                                                                               single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: linear
US-09-081-975-19
                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-277-660A-1
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US-08-424-957-20
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                                                                                                                 Sequence 4, Application US/08277660A
Patent No. 5702908
GENERAL Incordant Sequence 4
GENERAL Incordant Steven M.
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES. 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
GITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-424-957-1

| Sequence 1, Application US/08424957 |
| Sequence 1, Application US/08424957 |
| Patent No. 2770377 |
| Patent No. 2770377 |
| APPLICANT: Dicksley, Steven M. APPLICANT: Lane, David P. |
| TITLE OF INVENTION: Interruption of Binding of MDM2 and P53 |
| TITLE OF INVENTION: Interruption of Binding of MDM2 and P53 |
| TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50 |
| CORRESPONDENCE ADDRESS: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: Plehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 |
| CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94111-4187
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
ODFRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPK: 910 277299
INFORMATION FOR SEQ ID NO: 4:
SEQUINCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.
Matches 14; Conservative
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1 OETFSDLWKLLPEN 14
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                                                                            RESULT 10
US-08-277-660A-4
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COMPUTER: IN PC-OMPATION

CONFIDENTIANG SYTEM: PC-DOS/NB-DOS

SOFTAMENT APPLICATION NDATA:

PRIOR APPLICATION NDATA:

PRIOR APPLICATION NDATA:

PRIOR APPLICATION NDATA:

PRIOR APPLICATION NUMBER: USG04/24,957

CLASSITICATION NUMBER: USG04/24,957

CLASSITICATION NUMBER: USG04/24,957

CLASSITICATION NUMBER: USG04/24,957

CLASSITICATION NUMBER: USG04/24,957

PRIOR APPLICATION NUMBER: USG04/24,957

PREEMING NOW: SOLD IN D. IS BECOME TO BE USG04/24,957

TELEPHONE: (415) 788-1289

TELEPHONE: (415) 788-1289
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APPLICANT: Picksley, Steven M.
APPLICANT: David P.
TILLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
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                                                                                                                                                           Length 15;
                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco CITY: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentun Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                      Query Match 73.3%; Score 77; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.8e-06;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 1, Application US/09035686
; Patent No. 6153391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET UMBER: A-C
TELECOMMUNICATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Dreger, Walter H. REGISTRATION NUMBER: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RY: United States 94111-4187
                   LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
SEQUENCE CHARACTERISTICS:
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                                                                                         unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                       ;
TOPOLOGY:
US-08-424-957-20
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RESULT 14

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Sequence 20, Application US/09035686

APPLICANT: Pickaley, Steven M.

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSE: Flehr, Hobbach, Test, Albritton & Herbert

STREET: Rour Embarcadero Center, Suite 3400

STATE: California

COUNTRY: United States

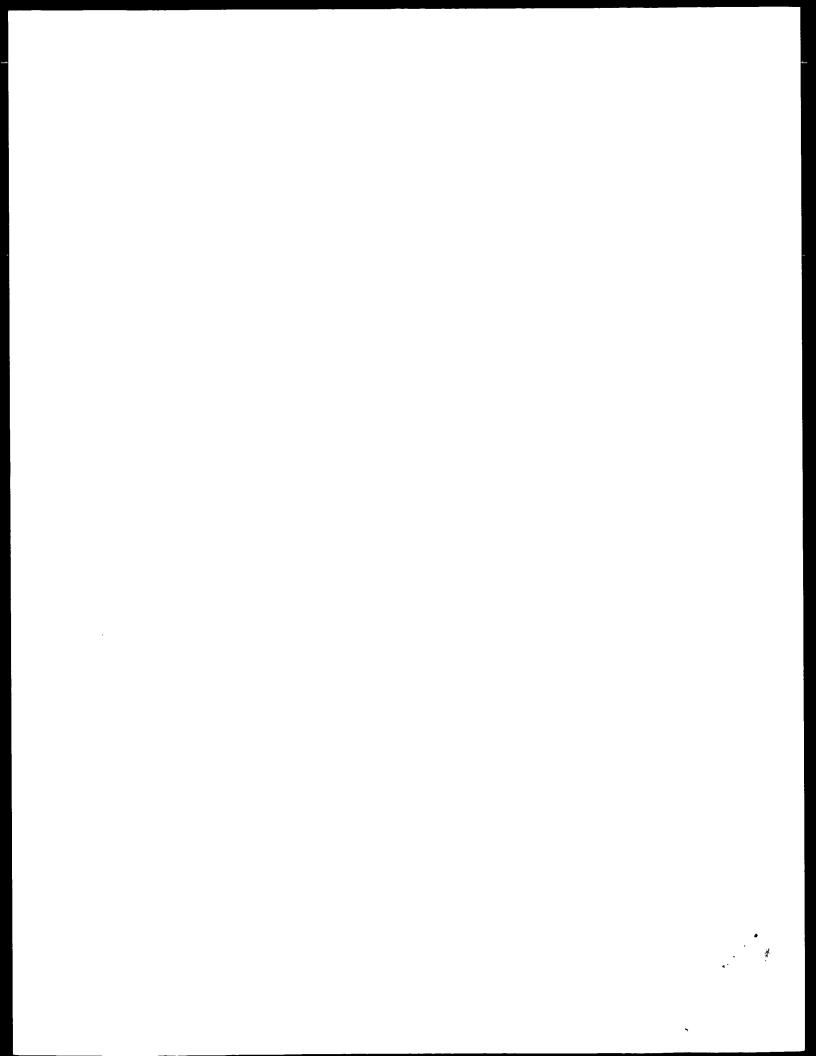
ZIP: 9411-4187
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APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Pherapeutic Application Thereof
NUMBER OF SEQUENCES: 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Flehr, Hobbach, Test, Albritton & Herbert
Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, 'CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                       PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 5, Application US/08277660A
Patent No. 5702908
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: (415, 22, TELEX: 910 277299; INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids TVPE: amino acid
                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: A-f
TELECOMMUNICATION INFORMATION:
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TELEPHONE: (415) , ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS: ADDRESS: Flehr, Ho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PPLSQETFSDLWKL 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 15
US-08-277-660A-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-035-686-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
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WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM. PC-DoS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATE:
PLOS/MS-DOS
FILING DATE: 20-JUL-1994
CLASSIFICATION NUMBER: US/08/277,660A

FILING DATE: 20-JUL-1994
CLASSIFICATION 514
ATTORNEY/AGENT INFORMATION:
NAME: ACCEST WHIRE: 24,19
REGISTRATION INFORMATION:
FELENCHMICATION INFORMATION:
FELENCHMICATION NUMBER: A.60244/WHD
FELENCHMICATION NUMBER: A.60249
FELENCHMICATION NUMBER: A.60244/WHD
FELENCHMICATION NUMBER: A.60249
FELENCHMICATION NUMBER: A.60244/WHD
FELENCHMICATION NUMBER: A.60249
FELENCHMICATION NUMBER: A.60244/WHD
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4: /cgn2 6/ptodata/1/paa/USO7 NEW COMB.pep:*

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7: /cgn2_6/ptodata/1/paa/USO8 NEW COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                                             OM protein - protein search, using sw model
                                                                                                                                                                                                                                                    US-09-403-440A-2
105
1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                            Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence 9, Appli	141		~	286	9	12,	7	53,	117	-	3	33,	35	. 9	ω,	127	• •	49,	28,	61,	3.	94	70.	70,	30,
SUMMARIES	ΙD	US-60-438-805-9	US-10-211-088-141	US-09-701-080C-22	US-09-701-080C-27	US-09-189-702A-286	US-10-049-428-6	US-09-701-080C-12	US-10-049-428-7	US-10-268-332-53	US-10-225-567A-1177	US-09-189-702A-7	US-09-281-717A-31	US-09-281-717A-33	US-09-281-717A-35	US-09-830-693B-6	US-09-830-693B-8	US-10-203-138A-12722	٥.	US-10-348-131-49	US-10-283-599-28	US-10-263-162-61	US-10-169-026-3	US-10-225-567A-942	PCT-US02-28371-70	US-10-236-878-70	US-09-552-802B-30
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& Query	Match	87.6	77.1	66.7	66.7	56.2	39.0	38.1	33.3	30.0	30.0	29.5	29.5	29.5	29.2	29.5	29.5	29.5	29.5	28.6	28.6	28.6	٠	27.6	27.6	27.6	27.6
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TYPE: PRT

Sequence 54, Appl	Sequence 6, Appli	Sequence 149, App	Sequence 11, Appl		36	297.	219, 7	9. Ap	60	1519	257			2904	67	18,	67	Sequence 463, App
US-10-319-786-54	US-10-190-866A-6	US-10-283-599-149	US-09-632-036E-11	US-60-438-805-3	US-10-174-717A-36	US-60-427-045-297	US-10-283-599-219	US-10-232-410-9	US-10-299-867-60	US-10-203-138A-15194	US-09-641-528-25751	US-09-641-528-29042	US-09-641-528A-25751	US-09-641-528A-29042	US-10-029-206A-67	US-10-301-644-18	US-10-028-075B-67	US-09-641-528-463
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27.6	9.17	27.6	27.1	27.1	26.7	26.7	26.7	26.7	26.7	26.7	25.7	25.7	25.7	25.7	25.7	25.7	25.7	25.7
53	23	01	28.5	28.5	28	28	28		28	28	27	27	27	27	27	27	27	27
27	87	29	30	31	32	33	34	32	36	37	38	39	40	41	42	43	44	45

# ALIGNMENTS

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US-10-211-088-141

Sequence 141, Application US/10211088
GENERAL INFORMATION

APPLICANT: Bright, Gary R.

APPLICANT: Premkumar, D. David

APPLICANT: Chen, Yih-Tai

TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding

FILE REPRESENCE: 01-1022-US

CURRENT APPLICATION NUMBER: US/10/211, 088

CURRENT FILING DATE: 2002-10-15

PRIOR APPLICATION NUMBER: 60/309,395

PRIOR APPLICATION NUMBER: 60/309,395

PRIOR FILING DATE: 2001-08-01

PRIOR FILING DATE: 2001-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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             Sequence 9, Application US/60438805;
GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS;
TITLE OF INVENTION: MIXED ELEMENT CAPTURE AGENTS;
FILE REFERENCE: UTSD:935USP1;
CURRENT APPLICATION NUMBER: US/60/438,805;
CURRENT FILING DATE: 2003-01-09;
NUMBER OF SEQ ID NOS: 12;
SOFTWARE: Patentin Ver. 2.1;
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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SOFTWARE: Patentin version 3.1
SEQ ID NO 141
LENGTH: 15
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US-60-438-805-9
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                                                                                                                                                                                                                                                                                   APPLICANT: Sidney, John
APPLICANT: Kast, W. Martin
APPLICANT: Southwood, Scott
APPLICANT: Spimmune, Inc.
ITLE OF INVENTION: HAA Binding Peptides and Their Uses:
ITLE OF ILLING HAA BINDER: US/09/189, 702A
CURRENT APPLICATION NUMBER: US/09/189, 702A
CURRENT FILING DATE: 1998-11-10
PRIOR PILING DATE: 1994-03-04
NUMBER OF ERQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 56.2%; Score 59; DB 5; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0046; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: CALCISE, Ian G.
APPLICANT: Xu, Weiming
APPLICANT: Xu, Weiming
APPLICANT: Iniu, Lizhi
TITLE CANT: Liu, Lizhi
FILE REPERENCE: HO-P02380USO
CURRENT APPLICATION WHERE: US/10/049,428
CURRENT FILING DATE: 2000-07-28
PRIOR APPLICATION WHERE: US/10/049,428
PRIOR PRILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 2000-06-30
NUMBER OF FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.1
Indels
0;
  0; Mismatches
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                                                                                                                                                                                                         Sequence 286, Application US/09189702A; GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
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Matches 9; Conservative
Matches 13; Conservative
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                                                  4 SQETFSDLWKLLP 16
                                                                                 1 SQETFSDLWKLLP 13
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US-09-189-702A-286
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US-10-049-428-6
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GENERAL INFORMATION
GENERAL INFORMATION
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION TRANSCRIPTIONAL REGULATION
TITLE OF INVENTION TRANSCRIPTIONAL REGULATION
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: GB 9811303.8
PRIOR PRILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: GB 9900157.0
PRIOR APPLICATION NUMBER: GB 9900157.0
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
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GENERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULER AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: US 9811303.8
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR FILING DATE: 1998-05-26
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
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                                                                                                                       77.1%; Score 81; DB 6; Length 15; 100.0%; Pred. No. 2e-06; tive 0; Mismatches 0; Indels
                                             , OTHER INFORMATION: Binding domain US-10-211-088-141
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          ORGANISM: Artificial sequence
                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 15; Conservative
                                                                                                                                                                                                                                                         1 PPLSQETFSDLWKLL 15
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Best Local Similarity
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US-09-701-080C-22
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TITLE OF INVENTION: IMMUNE- AND COLON- RELATED TISSUES
                          FILE REFERENCE: D0042A CIP
CURRENT APPLICATION NUMBER: US/10/268,332
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US. 60/235,713
PRIOR FILING DATE: 2000-09-27
PRIOR PLING DATE: 2001-01-16
PRIOR PLING DATE: 2001-01-16
PRIOR PLING DATE: 2001-01-17
PRIOR PLING DATE: 2001-01-17
PRIOR PLING DATE: 2001-07-13
PRIOR PLING DATE: 2001-08-17
PRIOR PLING DATE: 2001-08-17
PRIOR PELING DATE: 2001-08-17
PRIOR PLING DATE: 2001-09-26
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Matches 7; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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5 ESFSDELWK 13
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TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY3, EXPRESSED HIGHI
                                                                                            APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 I
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILLE REPERENCE: N134770 GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 1098-05-26
PRIOR PILING DATE: 1998-05-26
PRIOR FILING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PARCENT NOS: 36
SOFTWARE: PARCENT NOS: 36
SOFTWARE: PARCENT NOS: 36
SEQ ID NOS: 36
SEQ ID NOS: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 7, Application US/10049428
GENERAL INFORMATION:
APPLICANT: Charles, Ian G.
APPLICANT: Charles, Ian G.
APPLICANT: Liu, Lizhi
TITLE OF INVENTION: Unducible Screen for Drug Discovery
TITLE OF ILLNG DATE: 200-07-28
CURRENT APPLICATION NUMBER: US/10/049,428
CURRENT APPLICATION NUMBER: GB 9918077
PRIOR PILING DATE: 1999-07-30
PRIOR PILING DATE: 2000-06-30
PRIOR PILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     38.1%; Score 40; DB 5; Length 7; 100.0%; Pred. No. 3.5e+05; tive 0; Mismatches 0; Indels
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                                               Sequence 12, Application US/09701080C GENERAL INFORMATION:
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; Sequence 53, Application US/10268332
; GENERAL INFORWATION:
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ORGANISM: Artificial Sequence
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Matches 7; Conservative
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RESULT 7 /
US-09-701-080C-12
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Sequence 1177, Application US/10225567A

Sequence 1177, Application US/10225567A

GENERAL INFORMATION:
APPLICANT: LifeSpan Bloacciences
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS FILE REFERENCE: 1220-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT APPLICATION NUMBER: 60/257,144
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR PILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 1177
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30.0%; Score 31.5; DB 6; Length 14; 77.8%; Pred. No. 1.4e+02;
                                                         Indels
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APPLICANT: Kast, W. Martin
APPLICANT: Southwood, Scott
APPLICANT: Epimmune, Inc.
TITLE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30.0%; Score 31.5; DB 6; 53.3%; Pred. No. 1.6e+02; tive 2; Mismatches 2;
                                                            1; Mismatches
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Sequence 33, Application US/09281717A

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APPLICANT: Darimont, Beatrice
APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Fletterick, Robert
APPLICANT: Fletterick, Robert
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Nathons BINDING
TITLE OF INVENTION: COACTIVACE BINDING
FILE REFERENCE: 9811-008-999
CURRENT APPLICATION NUMBER: US/09/281,717A
CURRENT FILING DATE: 1998-03-30
PRIOR FILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
SEQ ID NO 31
LENGTH: 12
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                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                            Query Match 29.5%; Score 31; DB 5; Length 9; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                          ; OTHER INFORMATION: p53.24V9 peptide 1323.02
US-09-189-702A-7
CURRENT APPLICATION NUMBER: US/09/189,702A CURRENT FILING DATE: 1998-11-10 PRIOR APPLICATION NUMBER: US 08/205,713 PRIOR FILING DATE: 1994-03-04 WUMBER OF SEQ ID NOS: 380 SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: (5)..(5)
OTHER INFORMATION: Leu --> Arg (L454R)
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LOCATION: (7)...(7)
OTHER INFORMATION: Leu --> Arg (L456R)
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; OTHER INFORMATION: Glu -->Lys (E457K)
US-09-281-717A-31
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GENERAL INFORMATION:
APPLICANT: Baxter, John
                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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Best Local Similarity 60.0%
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US-09-281-717A-33
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APPLICANT: Darimont, Beatrice
APPLICANT: Parimont, Mediun
APPLICANT: Feng, Wediun
APPLICANT: Feng, Wediun
APPLICANT: Rushner, Peter
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wester
APPL
                          APPLICANT: Darimont, Beatrice
APPLICANT: Darimont, Beatrice
APPLICANT: Darimont, Beatrice
APPLICANT: Peng, Weijun
APPLICANT: Fetterick, Robert
APPLICANT: Weijun
APPLICANT: West, Brian
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Matches 6; Conserv
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GENERAL INFORMATION:
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### SPELICANT: Shidaw, Andrew
### APPLICANT: Shidaw, Andrew
### APPLICANT: Shidaw, Andrew
### APPLICANT: Shidaw Seeffery
#### APPLICANT: Creene, Geoffery
#### APPLICANT: Greene, Geoffery
#### TITLE OF INVENTION: ACTIVITY
#### TITLE OF INVENTION: ACTIVITY
### TITLE OF INVENTION: ACTIVITY
### FILE REPERBNCE: 9811-013-999
### CURRENT PELLING DATE: 1999-00-01-39
### PRIOR PAPLICATION NUMBER: US 60/079,956
### PRIOR APPLICATION NUMBER: US 60/079,956
### PRIOR PAPLICATION NUMBER: US 60/113,146
### PRIOR PAPLICATION NUMBER: US 60/113,146
### PRIOR PAPLICATION NUMBER: US 60/113,014
### PRIOR PAP
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1 PPLSQETFSDLWKLLPENG 19
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ORGANISM: Homo sapiens
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Best Local Similarity
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Sequence 74, Appl
Sequence 8, Appli
Sequence 10, Appl
Sequence 11, Appl
Sequence 11, Appl
                                                                                                                     February 12, 2003, 11:13:52; Search time 377.625 Seconds (without alignments) 32.439 Million cell updates/sec
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1: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

2: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

3: \cgn2 6/ptodata/1/paa/USO8 COMB.pep:*

4: \cgn2 6/ptodata/1/paa/USO8 COMB.pep:*

5: \cgn2 6/ptodata/1/paa/USO81 COMB.pep:*

7: \cgn2 6/ptodata/1/paa/USO82 COMB.pep:*

9: \cgn2 6/ptodata/1/paa/USO83 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO84 COMB.pep:*

11: \cgn2 6/ptodata/1/paa/USO85 COMB.pep:*

12: \cgn2 6/ptodata/1/paa/USO85 COMB.pep:*

11: \cgn2 6/ptodata/1/paa/USO85 COMB.pep:*

12: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

13: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

14: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

15: \cgn2 6/ptodata/1/paa/USO92 COMB.pep:*

16: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

17: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

18: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

20: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

21: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

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20: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

21: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

24: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*
GenCore version 5.1.3
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US-09-214-371-74
US-10-155-059-8
US-10-155-059-10
US-10-155-059-11
US-10-155-059-11
                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                         4569144 seqs, 644733110 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                   protein search, using sw model
                                                                                                                                                                                                                              105
1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                             US-09-403-440A-2
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Match Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.3
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                                                                                                                            Run on:
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Sequence 1109, Ap Sequence 1109, Ap Sequence 1109, Ap Sequence 16, Appli Sequence 2, Appli Sequence 3, Appli Sequence 2, Appli Sequence 2, Appli Sequence 5, Appli
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Sequence 13, Appl
Sequence 26, Appl
Sequence 4, Appli
Sequence 4, Appli
Sequence 14, Appli
Sequence 18, Appli
Sequence 6, Appli
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Appli
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Appli
                              Sequence 14, App.
                                              Sequence 15, App.
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Sequence 20, Appl
                                                                                         Sequence 13,
                                                                                                               Sequence 19,
                                                                                                                                                 Sequence 4, A
Sequence 3, A
Sequence 4, A
                                                             Sequence 73
                                                                                                                                   Sequence 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8,
Sequence 8,
                                                                           Sequence 7
                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 2, Application US/09403440A
GENERAL INFORMATION:
TITLE OF INVENTION: MATERIALS AND METHODS RELATING TO
TITLE OF INVENTION: INHIBITING THE INTERACTION OF p53 AND MDM2
FILE REFERENCE: MEMB25.001APC
CURRENT APPLICATION NUMBER: US/09/403,440A
CURRENT FILING DATE: 2000-01-19
FRIOR APPLICATION NUMBER: GF/GB98/01144
FRIOR APPLICATION NUMBER: GB 9708092.3
FRIOR PILING DATE: 1999-04-20
FRIOR FILING DATE: 1997-04-22
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FREUENCE FOR WINDOWS VERSION 4.0
                                                                                                                                                                                   US-09-458-297-1109
US-09-458-297-1109
US-09-458-297-1469
US-09-458-297A-1109
US-09-458-297A-1469
US-09-936-956-2
US-09-936-956-3
US-09-936-956-9
US-08-277-666-5
US-08-277-666-5
US-08-277-66-5
US-08-277
                          US-10-155-059-14

US-10-155-059-15

US-09-214-371-73

US-09-214-371-73

US-10-155-059-13

US-10-155-059-13

US-08-277-660-1

US-08-277-660-1

US-08-277-660-4

US-08-424-273-3

US-08-424-273-3
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US-09-936-956-13
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US-10-155-059-18
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US-08-398-139B-8
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US-08-132-284-4
US-08-398-139B-4
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   US-09-403-440A-2
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Gaps

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Indels

100.0%; Score 105; DB 18; 100.0%; Pred. No. 7.8e-09; iive 0; Mismatches 0;

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Query Match
Best Local Similarity
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US-10-155-059-10
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Sequence 8, Application US/10155059
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: AGelin, William
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: METHODS OF TREATMENT USING
ANTIBODIES
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.3%; Score 99; DB 16; Length 18; 100.0%; Pred. No. 6.3e-08; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
FLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                             APPLICANT: Pickšley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
IITLE OF INVENTION: Inhibitors of the
US-09-214-371-74; Sequence 74, Application US/09214371B; GENERAL INFORMATION:
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                                                                                                                                                                 Bottger, Volker
Bottger, Angelica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PPLSOETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100. Matches 18; Conservative
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                                                                                                                        Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE:
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LENGTH: 18
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                                                                                                                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
                                                                                                                             APPLICANT:
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Sequence 10, Application US/10155059
GENERAL INFORMATION:
APPLICANT: Kaelin, William
TITLE OF INVENTION: METHODS OF TREATMENT USING
ITTLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%; Score 99; DB 25; Length 20; 100.0%; Pred. No. 7.2e-08; tive 0; Mismatches 0; Indels
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OPERATING SYSTEM: Windows
OPERATING SYSTEM: Windows
SOFTWARE: PastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIPICATION: <understand number of the company of the
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APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
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SEQUENCE DESCRIPTION: SEQ ID NO: 10:
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 20 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 02110
COMPUTER READABLE FORM:
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Best Local Similarity 100.03
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boston
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NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 18, Application US/08244476; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                 MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: <URKNOWN>
INFORMATION FOR SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
    NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC COMPAT
OPERATING SYSTEM: PC-DC
SOFTWARE: PATENTIN Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.'
Matches 18; Conservative
                                                                                 CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U.S.A.
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US-10-155-059-16
is General 16, Application US/10155059
is GENERAL INFORMATION:
i APPLICANT: Kaelin, William
i APPLICANT: Kaelin, William
i TITLE OF INVENTION: METHODS OF TREATMENT USING
i TITLE OF INVENTION: MESTHODS OF TREATMENT USING
i ANTIBODIES
ANTIBODIES
                                                                                                                              RESULT 5
US-10-155-059-11
; Sequence 11, Application US/10155059
; GENERAL INFORMATION:
; APPLICANT: Kaelin, William
; JOSt, Christine
; TITLE OF INVENTION: METHODS OF TREATMENT USING
; TITLE AMD USES OF THE AND PROTEINS THERETO, AND USES OF THE ANTIBODIES

ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ô
        Gaps
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        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SCHWARE: FastSEQ for Windows Version 2.0b CURRENT APPLICATION DATA:
APPLICATION DATE: 24-May-2002
FILING DATE: 24-May-2002
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
RECEISTRATION NUMBER: 30,628
REFERENE/POCKET NUMBER: 30,628
        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS: ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 20 amino acids
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
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                                                                     2 PPLSQETFSDLWKLLPEN 19
                                                1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: <Unknown>
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Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-155-059-11
Matches
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APPLICANT: Schranz, Peter
APPLICANT: Schranz, Peter
APPLICANT: Volkmann, Martin
APPLICANT: Volkmann, Martin
APPLICANT: Tesmer, Claudia
APPLICANT: Klein, Ralf
TITLE OF INVENTION: METHOD OF DETECTING p53-SPECIFIC
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 30
CORRESPONDENCE ADDRESS:
ADDRESSE: Venable, Baetjer, Howard & Civiletti, LLP
STREET: 1201 New York Avenue, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%; Score 99; DB 25; Length 20; 100.0%; Pred. No. 7.2e-08; Live 0; Mismatches 0; Indels
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
CLASSIFICATION: 4UNROWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
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US-08-424-273-1

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TITLE OF INVENTION: P53 PROTEIN FRAGMENTS AND USE THEREOF FOR THITLE OF INVENTION: DETECTING AND MONITORING DISEASED CONDITIONS NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.3%; Score 99; DB 6; Length 25; 100.0%; Pred. No. 9.3e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Jacobson, Price, Holman & Stern, PLLC STREET: 400 Seventh Street, N.W. CITY: Washington
                                                     PULGATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: WO 94/08241
FILING DATE: 30-SEP-1993
FRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 42 32 823.3
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Thinen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REPRENENCE/DOCKET NUMBER: 18644-94281-1
TELEPHONE: 202-962-4810
TELEPHONE: 202-962-4810
TELEPATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC comparible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Word Perfect 5.1 (ASC III)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,273
FILING DATE: 28-APR-1995
FILING DATE: 28-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 9213110
FILING DATE: 2-NOV-1992
FILING DATE: 2-NOV-1992
FILING DATE: 2-NOV-1993
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH 25 amino acids
APPLICATION NUMBER: US/08/244,476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-424-273-1; Sequence 1, Application US/08424273; Sequence I, Application US/08424273; APPLICANT:
                FILING DATE: 15-SEP-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 PPLSQETFSDLWKLLPEN 21
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOFOLOGY: linear
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D.C.
USA
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RESULT >
US-10-155-059-14
US-10-155-059-14
; Sequence 14, Application US/10155059
; GENERAL INFORMATION:
; GENERAL INFORMATION:
; JOSE, Christine
; TITLE OF INVENTION: METHODS OF TREATMENT USING
; TITLE OF INVENTION: METHODS OF TREATMENT USING
                                               Gaps
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      Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
  94.3%; Score 99; DB 8; Le
100.0%; Pred. No. 9.3e-08;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «Inknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 14: US-10-155-059-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-155-059-15;
Sequence 15, Application US/10155059;
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: MA
COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Disket
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                 1 PPLSQETFSDLWKLLPEN 18
                                                                                                                      2 PPLSQETFSDLWKLLPEN 19
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                                          Conservative
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Best Local Similarity 94.45
Matches 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
Query Match
Best Local Similarity
Matches 18; Conserv
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RESULT 13
US-10-155-059-13
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US-09-732-384-7
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                                                                                FEATURE:
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•
JOST, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: General Control
APPLICANT: Furet, Pascal
ITILE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: INHIBIT CONTROL
CURRENT APPLICATION NUMBER: P2092/26
FRIOR APPLICATION NUMBER: P27/EP97/03549
FRIOR FILING DATE: 1999-03-26
FRIOR PILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLIASSIFICATION ATM.

PRIOR APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 06/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELEGOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFRANCE | 10-345-6054
TELEFRANCE | 10-345-6054
TELEFRANCE | 10-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                    ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
                                                                                                           NUMBER OF SEQUENCES: 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 94.49
Watches 17; Conservative
                                                                                                                                                                                                      CITY: Boston
                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-214-371-73
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NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE ANTIBODIES
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                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                  88.6%; Score 93; DB 16; Length 22; 100.0%; Pred. No. 6.9e-07; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87.6%; Score 92; DB 21; Length 19; 100.0%; Pred. No. 8.4e-07;
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COMPUTER: IBM Compatible
ODERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7. Application US/09732384; GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
TITLE OF INVENTION: Inhibition of p53 Degradation; TITLE OF INVENTION: Inhibition of p53 Degradation; TITLE OF INVENTION: Inhibition of p53 Degradation; TITLE OF INVENTION: 1000-12-07; CURRENT APPLICATION NUMBER: US/09/732,384; CURRENT FLING DATE: 1999-12-08; PRIOR FILING DATE: 1999-12-08; NUMBER OF SEQ ID NOS: 10; SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kaelin, William Jost, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. No. -
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ADDRESSE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 13, Application US/10155059 GENERAL INFORMATION:
                                                                                                                                                                       NAME/KEY: VARIANT
LOCATION: (1)
SOTHER INFORMATION: X = Biotin-Ser
US-09-214-371-73
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PPLSQETFSDLWKLLPE 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6 PPLSQETFSDLWKLLPE 22
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APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TILLE ON TOWNION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 85.7%; Score 90; DB 25; Length 19; Best Local Similarity 88.9%; Pred. No. 1.7e-06; Matches 16; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     84.8%; Score 89; DB 16; Length 19; 94.1%; Pred. No. 2.5e-06; tive 1; Mismatches 0; Indels
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «UNKNOWN»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1996
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                               NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09214371B GENERAL INFORMATION:
                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                               TELEX: <Unknown>
NATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lane, David
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
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Best Local Similarity 94.14
Matches 16; Conservative
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US-09-214-371-1
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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RESULT 15

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APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OOPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Search completed: February 12, 2003, 11:24:05 Job time : 378.625 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 19:
                                                                                                                                                                                                                                 ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                   ; Sequence 19, Application US/10155059 ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                   ZIP: 02110
COMPUTER READABLE FORM:
                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 PLSQETFSDLWKLLPEN 18
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Matches 16; Conservative
                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                             CITY: Boston
                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                         STATE: MA
US-10-155-059-19
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:15:02 ; Search time 12.6667 Seconds (without alignments) 38.323 Million cell updates/sec Run on:

105 1 PPLSQETFSDLWKLLPENG 19 US-09-403-440A-2 Perfect score:

Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

42882 Total number of hits satisfying chosen parameters: 140259 seqs, 25548876 residues

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Published Applications AA:\*

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14: /cgn2\_6/ptodata/1/pubpaa/USO0\_NEW\_PUB.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

	Description	Semience 74 April	ď	2	Sequence 11. Appl	16.	14	Segmence 15, April	Segmence 73. Appl	4			19	9	23	20.	8	12		¢	rddw '77 annanhae
	ΙD	US-09-214-371-74	US-10-155-059-8	US-10-155-059-10	US-10-155-059-11	US-10-155-059-16	US-10-155-059-14	US-10-155-059-15	US-09-214-371-73	US-09-732-384-7	US-10-155-059-13	US-09-214-371-1	US-10-155-059-19	US-09-732-384-6	US-10-155-059-23	US-10-155-059-20	US-10-155-059-18	US-09-214-371-17	US-09-950-69-6	115-10-155-059-22	77-000 001 01 00
		10	12	12	12	12	12	12	10	10	12	10	12	10	12	12	12	10	10	12	1
	Query Match Length DB	18	20	20	20	20	20	20	22	19	19	19	20	15	16	20	20	12	15	α	)
ф	Query Match	94.3	94.3	94.3	94.3	94.3	89.5	89.5	88.6	87.6	85.7	84.8	81.9	73.3	70.5	70.5	9.89	62.9	61.9	57.1	
	Score	66	66	66	66	66	94	94	93	92	90	89	86	77	74	74	72	99	65	9	
	Result No.	H	7	m	4	ß	9	7	œ	Φ	10	11	12	13	14	15	16	17	18	19	

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; Sequence 8, Application US/10155059 ; Patent No. US20020147173A1

RESULT 2 US-10-155-059-8

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10 US-09-214-371-39 0 US-10-024-123-7 10 US-09-214-371-7 10 US-09-214-371-40 10 US-09-214-371-40 10 US-09-214-371-25 10 US-09-214-371-25 10 US-09-214-371-25 10 US-09-214-371-33 10 US-09-214-371-34 12 US-10-155-059-17 10 US-09-13-371-83 10 US-09-13-371-83 10 US-09-13-371-83 10 US-09-732-384-4 12 US-10-155-059-1 12 US-09-732-384-4 12 US-09-732-384-4 12 US-09-732-384-4 13 US-09-734-371-37 10 US-09-734-371-37 10 US-09-734-371-37 10 US-09-734-371-37 10 US-09-734-371-37 10 US-09-734-371-37 10 US-09-734-371-37	10 US-09-214-371-36 12 US-10-155-059-2 9 US-09-840-277-66 10 US-09-920-552-110 10 US-09-214-371-13
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#### ALIGNMENTS

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Sequence 74, Application US/09214371B

Patent No. US20010018511A1

GENERAL INFORMATION:

APPLICANT: Lane, David

APPLICANT: Bottger, Volker

APPLICANT: Bottger, Stephen

APPLICANT: Cartick

APPLICANT: Carcia-Echeveria, Carlos

APPLICANT: Puret, Pascal

TILLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2

FILE REFERENCE: 4-20937/A/PCT

CURRENT PLING DATE: 1999-03-26

PRIOR APPLICATION NUMBER: US/09/214,371B

CURRENT PLING DATE: 1999-03-26

PRIOR PLING DATE: 1999-03-26

PRIOR ELING DATE: 1999-07-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 74
LENGTH: 18
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US-09-214-371-74
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COMPUTER: IEM COMPATIBLE
COMPUTER: IEM COMPATIBLE
SOFTWARE: FASTENG for Windows
SOFTWARE: FASTENG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1999
FILING DATE: 12-MAY-1997
FILING DATE: 12-MAY-1997
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COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
ODERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
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PILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                             NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPRAK: 617-345-6054
TELEFAX: 617-345-1300
TELEX: CURROWN:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
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CORRESPONDENCE ADDRESS: ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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; Sequence 11, Application US/10155059
; Patent No. US20020147173A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
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Jost, Christine
JITLE OF INVENTION: METHODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                           APPLICANT: Kaelin, William
JOSt, Christine
TITLE OF INVENTION: METHODIES OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                  CURRENT ITE: DESCRICE
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COPERATING SYSTEM: Windows
SOFTWARE: FESTSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «Unknown>
PRIOR APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Eisenetein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                              ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
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                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                   CITY: Boston
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            GENERAL INFORMATION:
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                            ö
                                                            Gaps
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Query Match 94.3%; Score 99; DB 12; Length 20; Best Local Similarity 100.0%; Pred. No. 6.2e-09; Matches 18; Conservative 0; Mismatches 0; Indels
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ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                             Gaps
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         94.3%; Score 99; DB 12; Length 20; 100.0%; Pred. No. 6.2e-09; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: METHODS OF TREATMENT USING NBS-1, ANTIBODIES AND PROTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY, AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
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ADDRESSEE: Nixon Peabody LLP
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CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                      ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                 ; Sequence 14, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 15, Application US/10155059; Patent No. US200201417173A1; GENEAL INFORMATION: APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                    APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: «Unknown»
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 27
                                                                                                                           2 PPLSQETFSDLWKLLPEN 19
                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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                                                    Matches 18; Conservative
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Best Local Similarity
Matches 17; Conserva
                               Best Local Similarity
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US-10-155-059-14
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           Query Match
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JOSt, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
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CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 66/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                      INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acids
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 16, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
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INFORMATION FOR SEQ ID NO: 16:
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STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 PPLSQETFSDLWKLLPEN 19
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ZIP: 02110
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US-10-155-059-16
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Gaps

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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                              FEATURE: OTHER INFORMATION: Description of Artificial Sequence:peptide NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 87.6%; Score 92; DB 10; Length 19; Best Local Similarity 100.0%; Pred. No. 6.6e-08; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                          Query Match 88.6%; Score 93; DB 10; Length 22; Best Local Similarity 100.0%; Pred. No. 5.5e-08; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 7, Application US/09732384
Fatent No. US2002012397A1
GENERAL INCRMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Yuan, Zhi-Min
TITLE OF INVERTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT APPLICATION NUMBER: 60/169,816
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
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ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 13, Application US/10155059
; Patent No. US20020147173A1
; GENERAL INFORMATION:
                                                                                                                                                                      LOCATION: (1)

COTHER INFORMATION: X = Biotin-Ser
US-09-214-371-73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
SEQ ID NO 73
LENGTH: 22
TYPE: PR
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 27
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ZIP: 02110
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                              1 PPLSQETFSDLWKLLPE 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-155-059-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
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  JOBL, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89.5%; Score 94; DB 12; Length 20; 94.4%; Pred. No. 3.5e-08; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: Windows
SOFTWARE: FEASTERQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION DAMER: US/09/081,975
APPLICATION NUMBER: 06/046,207
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                               ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
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SEQUENCE CHARACTERISTICS:
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
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Best Local Similarity 94.44
Matches 17; Conservative
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APPLICANT: Bottger, Angelic
APPLICANT: Pickeley, Steph
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heir
APPLICANT: Garcia-Echeverri
APPLICANT: Furet, Paecal
TITLE OF INVENTION: Inhibit
FILE REPERENCE: 4-20937/A/I
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                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                                                                            ZIP: 02110
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Gaps

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1 PLSQQTFSDLWKLLPEN 17
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ZIP: 02110
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Best Local Similarity
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US-10-155-059-19
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US-09-732-384-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 85.7%; Score 90; DB 12; Length 19; 88.9%; Pred. No. 1.3e-07;
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASTSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION ATA:
APPLICATION ATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/46,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                           NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
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Bottger, Volker
Bottger, Nagelica
Botsley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 13: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
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SEQ ID NO 1
LENGTH: 19
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Matches 16, Conservative
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US-09-214-371-1
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APPLICANT:
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NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Comparible
OOPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                      JOST, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/09732384
Patent No. US/20020132977A1
GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
TILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059

PILING DATE: 24 MAY-2002

CLASSIFICATION NUMBER: US/09/081,975

PILING DATE: 12-MAY-1998

APPLICATION NUMBER: 0/046,207

PILING DATE: 12-MAY-1997

ATTORNEY/ACENT INFORMATION:

REGISTRATION: NUMBER: 30,628

REGISTRATION NUMBER: 30,628

REGISTRATION NUMBER: 30,628

REFERENCE/DOCKET NUMBER: 30,628
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                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                             STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/169,816
                                                                                                                                                                               ANTIBODIES
; Sequence 19, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 amino acids
                                                                              APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                         NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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PRIOR FILING DATE: 1999-12-08

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APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 06/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
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SEQUENCE DESCRIPTION: SEQ ID NO: 20:
                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                               Sequence 20, Application US/10155059; Patent No. US20020147173A1; GENERAL INPORMATION:
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                                                                                                                                                                                                                                        APPLICANT: Kaelin, William
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 617-345-1300
TELEX: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 70.5'
Best Local Similarity 93.3'
Matches 14, Conservative
         2 PLSQETFSDLWKLLPE 17
                                                     1 PLSOETFEDLWSLLPD 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 PLSQETFSDLWKLLP 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                              US-10-155-059-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-155-059-20
                                                                                                                        RESULT 15
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Jost, Christine
JILE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                                                       0; Gaps
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                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Protein COTHER INFORMATION: fragment not in inhibitory p53 polypeptide US-09-732-384-6
                                                                                                                                                                                                                                                    Query Match 73.3%; Score 77; DB 10; Length 15; Best Local Similarity 100.0%; Pred. No. 9.4e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match

Query Match

Pest Local Similarity 81.2%; Pred. No. 2.8e-05;

Matches 13. Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT IRM Compatible
CORRUTER: IRM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION - CURROWN>
PRIOR PRELICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INCRMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 23, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 16 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Disket
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <Unknown>
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                                                                                                                                                                                                                                                                                                                                             5 QETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                      1 OETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Boston
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                                             SEQ ID NO 6
LENGTH: 15
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TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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0
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                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
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February 12, 2003, 11:10:52; Search time 47.5 Seconds (without alignments) 53.300 Million cell updates/sec
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                            OM protein - protein search, using sw model
                                                                                                                                                Run on:
```

Title: Perfect score: Sequence:	US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19
Scoring table:	BLOSUM62 Gapop 10.0 , Gapext 0.5
Searched:	908470 seqs, 133250620 residues
Total number of	Total number of hits satisfying chosen parameters: 320064
Minimum DB seq length: 0 Maximum DB seq length: 25	.ength: 0 .ength: 25

. 08	1 100%	45 summaries	
Match	latch	first	
ing: Minimum	Maximum M	Listing 1	
Post-processing:			

Database :		A Geneseq 101002:	01002:*		
		1: /SIDS2/g	cgdata/g	/SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:	DAT: *
		2: /SIDS2/g	cgdata/g	SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:	DAT: *
		3: /SIDS2/g	cgdata/g	'SIDS2/gcgdata/geneseq/genesecp-embl/AA1982.DAT:	DAT: *
		4: /SIDS2/g	cgdata/g	'SIDS2/gcgdata/geneseq/geneseqp-embl/AA1983.DAT:	DAT: *
		5: /SIDS2/g	cgdata/g	SIDS2/gcgdata/geneseq/genesegp-embl/AA1984.	DAT: *
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		7: /SIDS2/g	cgdata/g	'SIDS2/gcgdata/geneseq/genesegp-emb1/AA1986.DAT	DAT: *
		8: /SIDS2/g	cgdata/g	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:	DAT: *
		9: /SIDS2/g	cgdata/g	SIDS2/gcgdata/geneseq/geneseqp-embl/AA1988.DAT:	DAT: *
		10: /SIDS2/	gcgdata/	/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT	.DAT:*
		11: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1990.DAT:	.DAT: *
		12: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/genesegp-embl/AA1991.DAT:	.DAT: *
		13: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1992.DAT:	.DAT:*
		14: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-embl/AA1993.DAT:	.DAT: *
		15: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1994.DAT:	.DAT:*
		16: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1995.DAT:	.DAT: *
		17: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/genesegp-emb1/AA1996.DAT:	.DAT: *
		18: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1997.DAT:	.DAT: *
		`	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-embl/AA1998.DAT:	.DAT: *
	••	`.	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-embl/AA1999.DAT:	.DAT: *
		`.	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-embl/AA2000.DAT:	.DAT:*
		22: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2001.DAT:	.DAT: *
	• •	23: /SIDS2/	gcgdata/	SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2002.DAT:	.DAT: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		Description	D53 homologue TIP	p53 homologue TIP	p53 N-terminal pep	Immunodominant epi	Human p53 amino ac	p53/RB interaction	p53 biotinylated N	Human p53 peptide	Immunodominant epi	Immunodominant epi
SUMMARIES		O C	AAW82321	AAW82319	AAW37228	AAR54907	AAR51879	AAW47074	AAW37227	AAY06310	AAR54909	AAR54910
		DB	19	13	13	15	15	18	13	20	15	12
	Query	Length	19	19	18	25	25	19	22	15	15	15
df	Query	Match	100.0	100.0	94.3	94.3	94.3	92.4	88.6	77.1	73.3	73.3
		Score	105	105	66	66	66	97	93	81	77	77
	Result	No.	-	8	е	4	ស	9	7	80	σ	. 10

Disclosure, Fig 1; 52pp; English.

tein b #1. 3a mot bindi tide 9	ide #7. inylated ide #11. ide #4.		Human p53 W110-typ Human oncogenic pr Mdm/hdm antagonist Mdm/hdm antagonist Mdm/hdm antagonist Mdm/hdm antagonist Human oncogenic pr Promega protein ki	Target activatable p53 DR supermotif Miniature protein Human oncogenic pr Mdm/hdm antagonist Mdm/hdm antagonist Mdm/hdm antagonist Mdm/hdm antagonist Human oncogenic pr Human oncogenic pr
AAR89914 AAB29157 AAG89500 AAG89730 AAY45227 AAB29159	AAB29163 ABB05528 AAB29167 AAB29160 AAB29161	AAY99001 AAY57799 AAB29162 AAB29164 AAB29149 AAB29158	AAW31188 AAB17087 AAB17087 ABB73170 ABB73182 AAW37201 AAR83012	ABB0236 AABB0246 AAB8269 AAB3190 AAB17077 ABB7317089 ABB73184 AAB73184 AAB73203
22 22 22 20 21 21	21 21 21 21 21	222222	233333333333333333333333333333333333333	23 23 23 13 13 13
15 15 15 17 17	15 14 15 15	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	444444444444444444444444444444444444444	122 123 125 125 125 125 125 125 125 125 125 125
73.3 73.3 73.3 73.3 70.5		67.6 65.7 65.7 63.3 63.3		588.11.588
C C C C C C C C C C C C C C C C C C C	74 73 71 71	71 70 69 69 66.5	1 A & & & & & & & & & & & & & & & & & &	61 61 61 61 61 61
111111111111111111111111111111111111111	17 18 19 20 21	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	2	0.000 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4

# ALIGNMENTS

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p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                        New substance with a mdm2 binding domain and coupling partner -useful for stabilising in cells without an efficient mdm2-mediated degradation pathway
              AAW82321 standard; Peptide; 19 AA.
                                                                                                                                                      98WO-GB01140.
                                                                                                                                                                       97GB-0008089.
                                                 (first entry)
                                                                 p53 homologue TIP peptide.
                                                                                                                                                                                                                         WPI; 1998-609975/51.
                                                                                                                                                                                       (UYDU-) UNIV DUNDEE.
                                                                                                                    WO9847919-A1.
                                                                                                                                                     20-APR-1998;
                                                                                                                                                                       22-APR-1997;
                                                 22-FEB-1999
                                                                                                                                     29-OCT-1998.
                                                                                                   Synthetic.
                                 AAW82321;
                                                                                                                                                                                                         Lane DP;
      RESULT 1
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0

Gaps

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This sequence is a peptide homologue of a region of p53 which binds to mand. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.
This sequence is a peptide homologue of a region of p53 which binds to mama. This peptide is used in the construction of a movel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells on or overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 tunction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2
                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                   Length 19;
                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                              100.0%; Score 105; DB 19;
100.0%; Pred. No. 5.7e-10;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW82319 standard; Peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 1; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-GB01144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97GB-0008092.
                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                                                                                             1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       p53 homologue TIP peptide.
                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-609932/51.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19 AA;
                                                                                                                                                                                                                                                                         19 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9847525-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-OCT-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW82319;
                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lane DP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW82319
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This represents a p53 N-terminal peptide fragment used in an Elisa TIP assay for analysing the interaction between human oncogenic protein MDM2 can do p53. The invention provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated cappetosis in tumour cells comprising a wild-type p53 and non-elevated compay and to identify and design inhibitors of MDM2, p53 binding. They can also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia conteractions, especially tumours and viral infections. The peptides can interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and caccumulation in normal cells. The peptides which mimic the MDM2 binding content in p53, have a significantly greater blocking activity compared content in wild-type p53.
                                                                                                                                                                                                                                                                                                                                      MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa TIP assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Compounds binding to MDW2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Boettger V, Chene P, Furet P, Garcia-echeverria C;
Lane D, Picksley S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                    Indels
                                                                                                                                                                                                                                                                                                  p53 N-terminal peptide fragment for Blisa TIP assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 94.3%; Score 99; DB 19; L Local Similarity 100.0%; Pred. No. 4.9e-09; les 18; Conservative 0; Mismatches 0;
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0
Pred. No. 5.7e-10;
                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 34; 45pp; English.
                                                                                                                                                                                        AAW37228 standard; peptide; 18 AA.
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0
  100.08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97GB-0007041.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96GB-0014197.
                                                        1 PPLSQETFSDLWKLLPENG 19
                                                                                          1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                  20-JUL-1998 (first entry)
                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lane D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NOVS ) NOVARTIS AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-100996/09.
  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boettger A, B
Hochkeppel H,
                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9801467-A2
                    19;
                                                                                                                                                                                                                              AAW37228;
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                    Matches
                                                                                                                                                      RESULT 3
                                                                                                                                                                            AAW37228
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0

0; Gaps

Query Match

**AAR5490** RESULT

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Zentgraf H;

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p53/RB interaction inhibiting peptide 6 (residues 11-29 of human p53).
                                                                                                                                                                                                                                                                                                                                                           Antibodies specific for p53 are detected by binding to immobilised fragments of the p53 gene product containing the antibody-binding region. Preferred fragments contain amino acids 1-241, 40-349, 40-393, 66-241, 66-393, 237-349, 237-393 and esp. 9-33, 37-52 or 368-386. See AAR51872-R51881 for sequences of these fragments.
                                                                                                                                                                                                                                      Non-radioactive detection of p53 specific antibodies - by capture on immobilised p53 or its fragments, then reaction with labelled second antibody, for diagnosis of tumours and suitable for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying compounds that modulate interaction of p53 and Rb protein - or those that bind to Rb protein, used to induce apoptosis, specifically for treatment of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Retinoblastoma gene; RB; p53 protein; interaction; inhibitor; tumour; apoptosis; modulator; medicine; veterinary; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94.3%; Score 99; DB 15; I
100.0%; Pred. No. 7.1e-09;
ive 0; Mismatches 0;
                                                                                                                                                   Volkmann M,
                                                                                                                    (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM STIFTUNG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW47074 standard; peptide; 19 AA.
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                                                                                                                                                     Tessmer C,
                                                                                                                                                                                                                                                                                                                              Claim 11; Page 19; 35pp; German.
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                                                  93WO-EP02666
                                                                                  92DE-4232823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSQETFSDLWKLLPEN 18
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Les 18; Conservative
                                                                                                                                                     Schranz P,
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                                                                                                                                                                                       WPI; 1994-135732/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 AA;
                                                                                                                                                                                                         N-PSDB; AAQ62365.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                  30-SEP-1993;
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                                                                                  30-SEP-1992;
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                14-APR-1994.
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                                                                                                                                                                                                                                                                                               screening
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                                                                                                                                                                                                                      cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human nuclear phosphoprotein p53; tumour suppressor gene product; anti-oncogene; cancer; tumour; antibody binding region; epitope.
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Pred. No. 7.1e-09;
0; Mismatches 0; Indels
                                                                                                                                                                                       Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR51879 standard; Protein; 25 AA.
                                                                                AARS4907 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 4; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Soussi T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%; S
100.0%;
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1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lubin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-167463/20
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Best Local Similarity
Matches 18; Conserv
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AAR51879

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Gaps

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Length 25; 0; Indels

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This peptide fragment of p53 is an inhibitor of the interaction between a p53 protein and a retinoblastoma (RB) protein. This peptide corresponds to residues 11-29 of human p53 (AAW47079). The interaction between p53 and Rb is found to be critical for determining whether or not a cell enters apoptosis. Apoptosis is prevented if interaction occurs. The interaction is between regions 1-7 or 290-393 of p53 and region 379-928 of Rb. The invention provides methods to identify compounds able to modulate interaction or binding between p53 and Rb protein. The method comprises combining p53 and Rb, or their fragments, with a test compound and detecting interaction/binding between them. These inhibitory compounds are used in human or veterinary medicine to modulate p53 activity and processes, specifically for inducing apoptosis in tumor cells (possibly also in cells infected by virus), in vivo or in vitro. Expression of these modulators by gene therapy methods is also contemplated. Other activities that can be affected are transcription repression, d1 arrest, DNA repair, homologous recombination and 20-5-5- exomuclease activity. Modulation of interaction with Rb may also
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa P2 assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Garcia-echeverria C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         p53 biotinylated N-terminal peptide fragment for Elisa P2 assay.
                                                                                                                                                                                                                                                                                                                                                                                                        Score 97; DB 18; Length 19;
Pred. No. 1.1e-08;
1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Boettger A, Boettger V, Chene P,
Hochkeppel H, Lane D, Picksley S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, Page 34; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW37227 standard; peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                        92.4%;
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96GB-0014197.
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                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 94.44
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                                stabilise p53.
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05-JUL-1996;
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This p53 biotinylated N-terminal peptide fragment is used in an Elisa P2 assay for analysing the interaction between human oncogenic protein MDM2 and p53. The invention provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2 and to identify and design inhibitors of MDMM2, p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia interactions, especially tumours and viral infections. The peptides can be administered masally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and site in p53, have a significantly greater blocking activity compared with wild-type p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New p53 regulatory protein (RB18A) useful as, e.g. sources of probes and primers to detect the transcription rate and abundance of RB18A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wound healing; graft rejection; reperfusion injury;
yyocardial infaction; stroke; traumatic brain injury;
neurodegenerative disease; ischaemia; toxaemia; infection; AIDS;
hepatitis; breast cancer; ovarian cancer; colon cancer; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RB18A, p53 regulatory protein; apoptosis; neoplasia; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This synthetic peptide corresponds to amino acids 13-27 of human pags. It was used to raise anti-peptide p53.1 and anti-RB18A antibodies in rabbit. RB18A (see AAVG6309) is a novel p53 regulatory protein of the invention. The antibodies were used in
                                                                                                                                                                                                                                                                                                                                                                                              Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                            88.6%; Score 93; DB 19; L
100.0%; Pred. No. 5.7e-08;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPE 17
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                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 100.
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                                                                                                                                                                                                                                                                                                                                                           22 AA;
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                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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AAR54910 standard; peptide; 15 AA.
                                                               29-NOV-1994 (first entry)
                                 AAR54910;
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                                                                                                                                                              Gaps
                      and
   immunoscreenings that led to the isolation of a cDNA clone (see AAX59124) encoding human RB18A. The invention provides methods and compositions for the diagnostic and therapeutic applications of RB18A, in particular for the diagnosis, prevention or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
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                                                                                                                          77.1%; Score 81; DB 20; Length 15; 100.0%; Pred. No. 3.2e-06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            transformation-associated protein.
                                                                                                                                                                                                                                                                                                    AARS4909 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Legros Y, Lubin R, Soussi T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92FR-0013110.
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                                                                                                                                                                                                                                                                                                                                                                (first entry)
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nes 14; Conservative
                                                                                                                                               Local Similarity 100.
nes 15; Conservative
                                                                                                                                                                                             2 PLSQETFSDLWKLLP 16
                                                                                                                                                                                                                            1 PLSOETFSDLWKLLP 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EURO-) LAB EUROBIO SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2 PPLSQETFSDLWKL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1994-167463/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 AA;
                                                                                                     15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                  29-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-MAY-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conditions.
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                                                                                                     Sequence
                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         states
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                                                                                                                                                                                                                                                                       RESULT 9
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AAR54910

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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-1953 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AARS4907-RS4921) are useful for detecting and monitoring cancerous and precancerous
                              cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                - for detecting and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
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                                                                                                                                                                                                                                                                                                                                                                                                                              New immuno:dominant epitope(s) of protein p53 - for detectir monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73.3%; Score 77; DB 15; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p53 protein binding motif corresp. to residues 16-30.
Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR89914 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                        Legros Y, Lubin R, Soussi T;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9602642-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-SEP-1996
                                                                                                                         Homo sapiens,
                                                                                                                                                                                                                                       02-NOV-1993;
                                                                                                                                                                                                                                                                            32-NOV-1992;
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                                                                                                                                                                                                11-MAY-1994.
                                                                                                                                                            WO9410306-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          conditions.
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p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                          New peptides of the invention which interfere with the binding of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                        New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fork head associated; FHA; domain; transcriptional control; DNA replication; DNA repair; cell cycle control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      / Match 73.3%; Score 77; DB 17; Length 15; Local Similarity 100.0%; Pred. No. 1.4e-05; les 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 1; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB29157 standard; Peptide; 15 AA.
19-APR-1995; 95US-0424957.
                          94US-0277660.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 OFTESDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 QETFSDLWKLLPEN 14
                                                                                                                                  Lane DP, Picksley SM;
                                                                           (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                         WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15 AA;
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                          20-JUL-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide

WPI; 2000-664872/64.

Disclosure, Fig 2; 92pp; English.

binding -

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The present invention describes isolated prepared P53 epitopes (I). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (1); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids, (3) a vaccine composition comprising (II), a unit dose of a peptide with less sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) am isolated nucleic acid encoding (I); and (5) am isolated nucleic acid encoding (II). (I) as cytostatic activity and can be used in vaccines. The vaccine composition is useful for treating or preventing cancer. (I) and (II) are useful as diagnostic agents and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be
The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL; cytotoxic T lymphocyte; T cell; immune response; tumour; cancer; vaccine; epitope; cytostatic.
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                                                                                                                                                                                                                                                                                                            .,
                                                                                                                                                                                                                                                                73.3%; Score 77; DB 21; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Southwood S, Chesnut R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        p53 DR 3a motif binding peptide exemplary sequence #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 113; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAG89500 standard; Peptide; 15 AA.
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                                                                                                                                                                                                                                                                                   Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                            1 PPLSOETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                  2 PPLSOETFSDLWKL 15
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                                                                                                                                                                                                                            15 AA;
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                                                                                                                                                                                                                            Sequence
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Keogh E;
                                                                                                                                                                                                                                                                Query Match
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The present invention describes isolated prepared P53 epitopes (I). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (I); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids; (3) a vaccine composition comprising (II), a unit dose of a peptide with less than 50 contiguous amino acids with 100% identity to the native peptide sequence of p53, and a pharmaceutical exciptent; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) are useful as diagnostic agents and for evaluating immune responses. Composition is useful for treating or preventing immune responses. Composition is bytopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine composition of (I). The ability to combine selected epitopes and
                composition of (I). The ability to combine selected epitopes and further, to modify the composition of the epitopes enhances the immunogenicity. The possible pathological side effects caused by infectious agents or whole pathological antigens, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                            Gaps
present in whole antigens can be avoided with the use of the vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL; cytotoxic T lymphocyte; T cell; immune response; tumour; cancer; vaccine; epitope; cytostatic.
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                                                                                                                                                                                                                    73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                           AAG89730 standard; Peptide; 15 AA.
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Keogh E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-2000; 2000WO-US33629
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p53 DR3 binding peptide #1.
                                                                                                                                                                                                                                                            14; Conservative
                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                       2 PPLSQETFSDLWKL 15
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                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                           present invention
                                                                                                                                                                              15 AA;
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Synthetic.
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                                                                                                                                                                                 Sequence
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            immunogenicity. The possible pathological side effects caused by infectious agents or whole protein antigens, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                       Immunisation; immunogenicity enhancer; rabies virus strain ERA; immunodominant T-helper cell determinant; nucleoprotein; antibody; post-translational modification; p53; cancer; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Methods of detecting post-translational modification level of p53, useful for diagnosis of cancer
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further, to modify the composition of the epitopes enhances the
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                                                                                                                                  73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 45; 59pp; English.
                                                                                                                                                                                                                                                                                                    AAY45227 standard; peptide; 17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Thurin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US04653.
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Best Local Similarity
                                                                             present invention.
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                                                                                                         15 AA;
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Search completed: February 12, 2003, 11:13:44 Job time: 47.5 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:12:02; Search time 37.2083 Seconds (without alignments) 105.216 Million cell updates/sec Run on:

US-09-403-440A-2

105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

**BLOSUM62** Scoring table:

Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters: 671580 seqs, 206047115 residues Searched:

9297

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SPTREMBL 21:\* Database :

sp\_arches:\*
sp\_bacteria:\*
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sp\_human:\*
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sp\_bacteriap:\*  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

sp\_archeap:\*

#### SUMMARIES

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Result		Query				
No.	Score	Match	Match Length DB	DB	ID	Description
-	77	73.3	25	4	Q9NP68	Ogno68 homo sanien
7	45	42.9	22	13	O9W6D7	Clap autlap 759w60
e	33	31.4	24	13	902J9	Ogorie and Street
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2	32	30.5	19	ß	P83003	PR3003 entamoeks k
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12	29	27.6	17	80	Q9XNQ1	O9xngl boophilus m
13	29	27.6	24	Ŋ	Q95V74	095v74 mvtilus dal
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O9nlc9 bos taurus O9tup6 sus scrofa O53422 xanthobacte O9uel0 homo sapien O9r3j2 ureaplasma P82326 pisum sativ O9qvs7 mus sp. lac O9ujz3 homo sapien O95mk2 papio cynoc		Q9t1d2 bostrychia Q64831 human adeno Q14347 homo sapien Q83960 influenzavi Q9f0p3 paracoccus P82674 bos taurus O84129 influenzavi	022064 dioscorea t Q83965 influenzavi Q9750 thermotoga Q95mK6 eulemur cor Q95mK5 varecia var Q95mK4 cheirogaleu
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#### ALIGNMENTS

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Gaps
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Bukaryota: Metazoa, Chordata, Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LA Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
Joyce K.A., Salisbury J., Sibson D.R., Vergote I., Walker C.;
Submitted (NOV-1999) to the BMEL/GenBank/DDBJ databases.
REMBL; AF209136; AAF3635.1; -.
REMBL; AF209129; AAF36355.1; -.
REMBL; AF209131; AAF36355.1; -.
REMBL; AF209131; AAF36357.1; -.
REMBL; AF209131; AAF36357.1; -.
REMBL; AF209131; AAF36350.1; -.
REMBL; AF209135; AAF36350.1; -.
REMBL; AF209135; AAF36360.1; -.
REMBL; AF30910910; AAF30910910; AAF3091091
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                                                                                                                                            01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                                                              PRT;
                                                                                                                                                                                                                                                                 Tumor suppressor p53 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 73.3
Best Local Similarity 100.
Matches 14; Conservative
                                                                          PRELIMINARY;
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RESULT 1
Q9NP68
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MEDLINE=99067015; PubMed=9851690;
Grechez-Cassiau A., Bernard M., Ladjali K., Rodriguez I.R., Voisin P.;
"Structural analysis of the chicken hydroxyindole-O-methyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              dallus allus (Chicken).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Archosauria, Aves, Neognathae, Galliformes, Phasianidae, Phasianinae,
                                                                                                                                    Gallus gallus (Chicken).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Archosauria, Aves, Neognathae, Galliformes, Phasianidae, Phasianinae,
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MEDLINE=84144799; PubMed=6322173;
Westaway D., Payne G., Varmus H.E.;
"Proviral deletions and oncogene base-substitutions in insertionally mutagenized c-myc alleles may contribute to the progression of avian bursal tumors.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                              / Match 42.9%; Score 45; DB 13; Length 22; Local Similarity 58.3%; Pred. No. 2.5; Losservative 4; Mismatches 1; Indels
                                                              01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NAR-2002 (TrEMBLrel. 20, Last annotation update)
Hydroxyindole-O-methyltransferase isoform A (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-DEC-2001 (TrEMBLrel. 19, Created)
01-BEC-2001 (TrEMBLrel. 19, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
C-myc oncogene, aa 44-62 from c-myc/LL3 (ALV provirus)
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EMBL; K01440; AAA48964.1; -.
InterPro; IPR002418; TF_Myc.
Pfam; PF01056; Myc_N_term; 1.
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                                 PRT;
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                                                                                                                                                                                                                                                                                                                                               EMBL; AF116455; AAD23444.1; -.
Methyltransferase; Transferase.
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                                 PRELIMINARY;
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-!- FUNCTION: THIS PROTEIN BINDS CALCIUM (BY SIMILARITY).
-!- SINGELLULUAR LOCATION: BNDOPLASMIC RETCULUM LUMEN.
-!- SINGELLULUAR LOCATION: BNDOPLASMIC RETCULUM LUMEN.
-!- SINGELLULUAR LOCATION: TROPHOZOITES.
-!- SIMILARITY: BELONGS TO THE CALRETICULIN FAMILY.
--- PROSITE; PSO0804; CALRETICULIN 1; PARTIAL.
--- PROSITE; PSO0804; CALRETICULIN 2; PARTIAL.
--- PROSITE; PSO0804; CALRETICULIN 2; PARTIAL.
--- PROSITE; PSO0804; CALRETICULIN 2; PARTIAL.
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                                                                                                                                                                                                                                      MEDLINE-84144799; PubMed-6322173; Westeraway D., Payne G., Varmus H.E.; Weoteraway D., Payne G., Varmus H.E.; "Proviral deletions and oncogene base-substitutions in insertionally mutagenized c_myc alleles may contribute to the progression of avian
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                              01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 21, Last annotation update)
RAV-2 proviral DNA, c-myc (Aa 44-67), allele LL3 (Fragment).
NAV-2 provirus.
Viruses, Retroid viruses; Retroviridae; Alpharetrovirus.
NCBI_TaxID=11864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31.4%; Score 33; DB 15; Length 24; 47.4%; Pred. No. 2.4e+02; tive 1; Mismatches 3; Indels
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01-0CT-2001 (TrEMBLrel. 18, Last sequence update)
01-0CT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                            Acad. Sci. U.S.A. 81:843-847(1984).
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Entamoeba histolytica.
Bukaryota, Entamoebidae, Entamoeba.
NCBI_TaxID=5759;
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                                                                                                                                                                                                                                                                                                                                              Proc. Natl. Acad. Sci. U.S.A. 8
EMBL; K01439; AAA51600.1; -.
InterPro; IPR002418; TF Myc.
Pfam; PF01056; Myc. N. term; 1.
NON TER 1 24 24
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                 bursal tumors."
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Q85618
Q85618;
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SEQUENCE
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                                                                                                                                                                                 STRAIN=HW383;
MEDLINE=20248365; PubMed=10789503;
Chida T., Okamura N., Ohtani K., Yoshida Y., Arakawa E., Watanabe H.;
"The complete DNA sequence of the O antigen gene region of Plesiomonas shigelloides serotype 017 which is identical to Shigella sonnei form
                                                                                                                                                                                                                                                                                                                  Houng H.H., Venkatesan M.M., "Genetic analysis of Shigella sonnei form I antigen: identification of a novel IS630 as an essential element for the form I antigen
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Acta Theriol. (Warsz) 45:143-146(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sorex araneus (Eurasian common shrew) (European shrew).
Wakaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Insectivora; Soricidae; Soricinae; Sorex.
                                                                                                            Shigella sonnei.
Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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Pred. No. 3.7e+02;
1; Mismatches 7; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                  25 AA; 2726 MW; A6CC5D24211CDAAD CRC64;
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                                                                            Last sequence update)
Last annotation update)
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01-UNN-2001 (TrEMBLrel. 17, Last sequence update)
01-UUN-2001 (TrEMBLrel. 17, Last_annotation update)
                                            25 AA
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                                                                                                                                                                                                                                                        Microbiol. Immunol. 44:161-172(2000).
                                                                   Created)
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EMBL; AB028134; BAA85070.1; -.
NON_TER 25
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EMBL; AF314827; AAK13419.1; -.
                                                   Q9SOUG;
01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-DEC-2001 (TrEMBLrel. 19,
ORF10S (Fragment).
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les 6; Conservative
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                                           PRELIMINARY;
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                                                                                                                                               NCBI_TaxID=624;
                                                                                                                                                                                                                                                                                                                                                       expression.";
                                                                                                                                                                                                                                                                                             STRAIN=HW383;
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SEQUENCE
                                                                                                                                      Shigella
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Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bos taurus (Bovine).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Cetartiodactyla, Ruminantia, Pecora, Bovoidea,
Bovidae, Bovinae, Bos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=99175163; PubMed=10075683; Jiang H., Okamura C.S., Lucy M.C.; Jiang H., Okamura C.S., Lucy M.C.; Isolation and characterization of a novel promoter for the bovine growth hormone receptor gene. 3790(1999).
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"The catalytic subunits of Ser/Thr protein phosphatases from
Caenorhabditis elegans: A biochemical and molecular biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: A PHOSPHOPROTEIN + H(2)0 = A PROTEIN PHOSPHATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      -!- SIMILARITY: BELONGS TO THE PPP FAMILY OF PHOSPHATASES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2BF73DBBB5F20359 CRC64;
                                                                                                                                               01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   077538; Q9TQV9;
01-NOV-1998 (TrEMBLrel. 08, Created)
01-WAR-2001 (TrEMBLrel. 16, Last sequence update)
01-WAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                PP1-like Ser/Thr protein phosphatase (Fragment). Caenorhabditis elegans.
                                                                                                           24 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL, Z77740; CAB01299.1; -.
HSSP; P08129; 1F7M
InterPro; 1PR004844; S/T_phosphtse.
ProDom; PD000252; S/T_phosphtse; 1.
Hydrolase; Iron; Manganese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Growth hormone receptor (Fragment).
                                                                                                           PRT;
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                                                                                                         PRELIMINARY;
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STRAIN=BREED HOLSTEIN;
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4 PPLFLEVFED 13
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Best Local Similarity 58.3
Matches 7; Conservative
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                                        NCBI_TaxID=6239;
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SEQUENCE
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Q95V74;
                                                                                                                survey."
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                      Jiang H., Lucy M.C.; "Identification of 5' untranslated variants of bovine growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
NCBI TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Orlovsky I.V., Sverdlova P.S., Rubtsov P.M.; "Microsatellites within the human growth hormone receptor gene."; Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ295613; CAC06615.1; -...
                                                                                     Jiang H., Lucy M.C.;
"Bovine GH receptor 5' UTR variants.";
"Bovine GH receptor 5' UTR variants.";
"Bobmitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF036297; AAC33310.1;
EMBL; AF036291; AAC33308.1;
EMBL; AF036293; AAC33310.1;
EMBL; AF036293; AAC33311.1;
EMBL; AF036295; AAC33311.1;
EMBL; AF036295; AAC33311.1;
EMBL; AF036295; AAC33314.1;
EMBL; AF036295; AAC33314.1;
EMBL; AF036295; AAC33314.1;
EMBL; AF036295; AAC33314.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 28.6%; Score 30; DB 4; Length 23; Local Similarity 83.3%; Pred. No. 7.2e+02; les 5; Conservative 1; Mismatches 0; Indels
                                                       Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases.
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SEQUENCE 23 AA; 2426 MW; 9AD833A0F1DD699E CRC64;
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01-FBB-1997 (TrEMBLrel. 02, Created)
01-FBB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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01-WAR-2001 (TrEMBLrel. 16, Last sequence update)
01-WAR-2001 (TrEMBLrel. 16, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                      23 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Growth hormone receptor (Fragment).
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2 DLWQLL 7
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2 DLWQLL 7
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NON TER
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
PPI-like Ser/Thr protein phosphatase (Fragment).
Caenorhabditis elegans.
Eukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
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Parasitiformes, Ixodida, Ixodidae, Boophilus.
                                                                                                                                                                                                                 Zeke T., Gergely P., Dombradi V.;
"The catalytic subunits of Ser/Thr protein phosphatases from Caenorhabditis elegans: A biochemical and molecular biological
                                                                                                                                                                                                                                                                                                                                  Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases. 
 -!- CATALYTIC ACTIVITY: A PHOSPHOPROTEIN + H(2)O=A PROTEIN +
                                                                                                                                                                                                                                                                                                                                                                                           PHOSPHATE.
-1- SIMILARITY: BELONGS TO THE PPP FAMILY OF PHOSPHATASES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4; Indels
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NON TER 17 17
SEQUENCE 17 AA; 1988 MW; 319F2D4DA7DA11F3 CRC64;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequ
01-NOV-1999 (TrEMBLrel. 12, Last anno
                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; Z77728; CAB61287.1; -.
HSSP; P08129; 1FJM.
INTERPRO; IPR004844; S/T_phosphtse.
ProDom; PD000252; S/T_phosphtse; 1.
Hydrolase; Iron; Manganese.
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Gaps
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Citrus sinensis (Sweet orange).
Citrus sinensis (Sweet orange).
Subaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
NCBI_TaxID=2711;
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                                                                                                                                           SEQUENCE FROM N.A.
Bierne N., Daguin C., Borsa P., Jollivet D., Viard F., Bonhomme F..
David P.;
                                             Elongation factor 1 alpha (Fragment).
Mytilus galloprovincialis (Mediterranean mussel).
Eukaryota; Metazoa; Mollusca; Bivalvia; Pteriomorphia; Mytiloidea; Mytilidae; Mytilus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Back J.-M., Kenerley C.M., Inthe arg2 gene of Trichoderma virens: Cloning and development of a
                                                                                                                                                                                          "Introgression patterns in the Mytilus edulis / Mytilus
galloprovincialis mosaic hybrid zone along the Atlantic coasts of
                                                                                                                                                                                                                                                                                                                                                                               ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27.1%; Score 28.5; DB 3; Length 24; 53.8%; Pred. No. 1.38+03; Live 2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                        27.6%; Score 29; DB 5; Length 24; 38.5%; Pred. No. 1.1e+03; rative 3; Mismatches 5; Indels
                                                                                                                                                                                                                                    Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          homologous transformation system.";
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases
EMBL; AF001029; AAB58298.1;
SEQUENCE 24 AA; 2782 MW; 8FFD15C27EAD244C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-1997 (TrEMBLrel. 04, Created)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Carbamoyl phosphate synthetase, small subunit (arg2).
Trichoderma virens.
                                                                                                                                                                                                                                                                                                       SEQUENCE 24 AA; 2875 MW; 1D0DEAD3FBD13DCB CRC64;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hypocreales, Hypocreaceae, Hypocrea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                         EMBL; AF424742; AAL24803.1; -.
                                                                                                                                                                                                                                                                                                                                                                        5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 53.8
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                   4 PPYSEARFMEIQK 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 SQETFSD-LWKLL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 SODYLSDHLWRAL 22
                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                             France."
                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P87182
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P87182
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RN [1]

RP SEQUENCE FROM N.A.

RA Liu C., Xu C., Lin B.;

RT "Molecular cloning of APETALA1 (AP1) gene from orange (Citrus r sinensis).",

RT sinensis).",

RL Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.

RL Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.

PT NON TER 20 2 2

SQ SEQÜENCE 20 As; 2493 MW; 3F1DOC6F8F5C1EA5 CRC64;

Query Match

Query Match

Acches 6; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

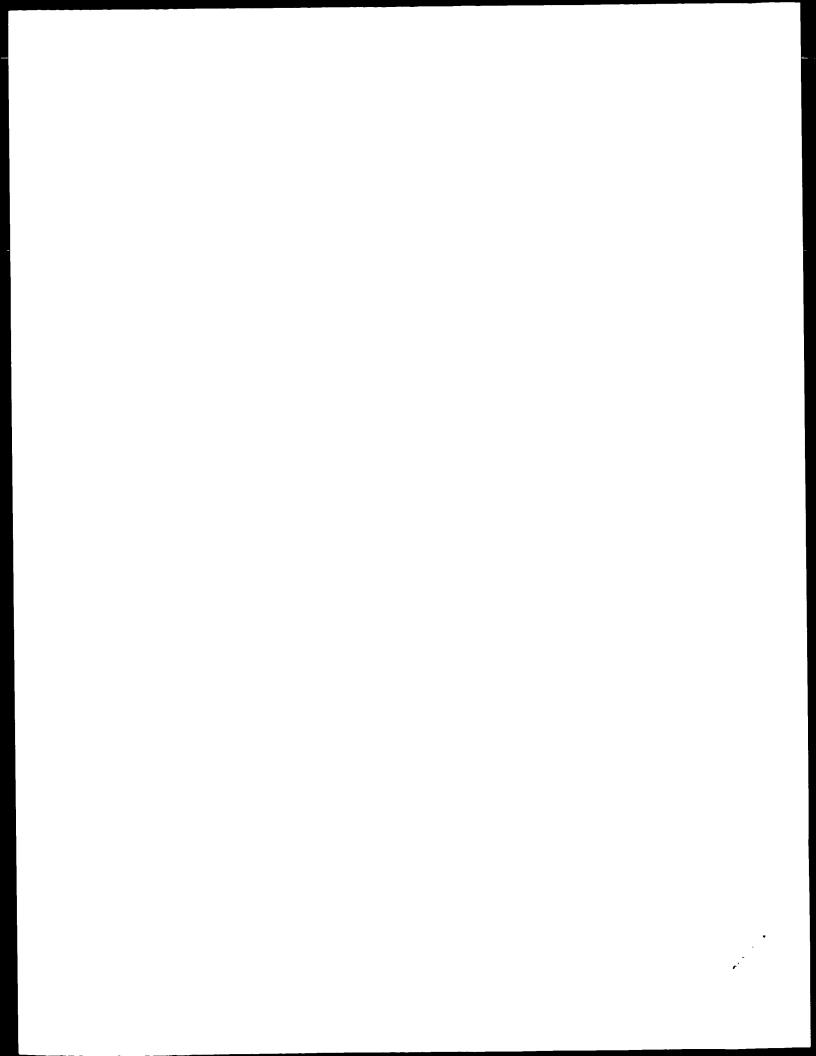
RATCHES 6; Conservative 2; Mismatches 3; Indels 0; Gaps O;

QY 3 LSQETFSDLWK 13

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Db 9 LTHESISDLOK 19

Search completed: February 12, 2003, 11:14:59
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

February 12, 2003, 11:11:31 ; Search time 10.2917 Seconds (without alignments) 76.572 Million cell updates/sec

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 segs, 41476328 residues Searched:

1520 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

SwissProt\_40:\* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## STIMMARTES

	cription	P22702 neurospora	P36231 iris hollan		P16353 heliothis z	•	spinacia				P80178 lymnaea sta		P80181 lymnaea sta		P24854 sus scrofa	P81490 phyllomedus		P04549 periplaneta			ın	σ	Q9ur66 candida alb					P82697 periplaneta	P82419 pachycondyl	P81217 equus cabal	LO.	0.0	93	39 lito
SUMMARIES	ID	Y3KD NEUCR	LECB_IRIHO	CRTC_SPIOL	HTF HELZE	CP35 PAPSP				LPCP_YEAST	NP1 LYMST	NP2_LYMST	NP4 LYMST	NP5_LYMST	IBP4 PIG	DMS6_PHYBI	UR2_SCYCA	HTF2_PERAM	HTF2_CARMO	URE1_MORMO	ACT6_SOYBN	CPAX_BOVIN	NDK_CANAL	AKH_TABAT	HTF1_PERAM	HTF_TABAT	NP3_LYMST							CR32_LITCE
	DB	ч	ч	Н	-1	н		-	-	-	Н	Н	н	-	н	-	-	Н	-	-	Н	Н	ч	<b>,-</b> -	<b>-</b>	-	-	Н	-	-	-		-	7
	Length	24	20	20	10	20	23	23	10	25	13	13	13	13	16	24	12	æ	10	12	17	18	21	ω	ω ,	10	13	17	18	19	19	19	20	22
916							22.9					21.0	21.0	21.0	21.0	21.0	20.5	20.0	20.0	20.0	20.0	20.0	20.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0	19.0
	Score	28.5	27	26	24	24		24		22.5	22	22	22	22	22	22	21.5	21	21	21	21	21	21	50	20	20	50	20	20	20	20	50	20	20
Result	No.	7	7	m	4	S	9	7	80	6	10	11	12	13	14	15	16	17	18	13	20	21	22	23	47.0	52	56	27	28	53	30	31	32	33

LECB IRIHO

ID LECB IRIHO

AC P36231:
DT 01-JUN-1994 (Rel. 29, Last sequence update)

DT 01-JUN-1994 (Rel. 29, Last annotation update)

DT 01-JUN-1994 (Rel. 29, Last annotation update)

DT 01-JUN-1994 (Rel. 29, Last annotation update)

DF 01-JUN-1994 (Rel. 29, Last annotation update)

EN acctyl-D-galactosamine-binding lectin subunit B (A-disaccharide-binding lectin subunit B) (Fragment).

RESULT 2

P56240 litoria cae	P24853 sus scrofa	P12664 oryctolagus	P19662 lactobacill	P21983 treponema p	P82423 pachycondyl	P82424 pachycondyl	P14086 gryllus bim	P80488 thiobacillu	P25419 tenebrio mo		
CR33_LITCE	IBP2_PIG	IRBP_RABIT	UDP LACCA	FLAA TREPH	PCW1_PACGO	PCW2_PACGO	AKHG GRYBI	CLP THICU	HTF TENMO	HTF NAUCI	URA7_HUMAN
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19.0	19.0	19.0	19.0	19.0	19.0	19.0	18.1	18.1	18.1	18.1	18.1
20	20	20	20	20	20	20	19	19	19	19	19
34	32	36	37	38	39	40	41	42	43	44	45

### ALIGNMENTS

e ooral	Auropean Boonnomatics Institutions as long as its content is in no way lied and this statement is not removed. Usage by and for commercial ites requires a license agreement (See http://www.isb-sib.ch/announce/nd an email to license@isb-sib.ch).
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etes ion N OF h a .	tric nd is sib is ib
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Sordariomycetes; Sordariomycetes; and expression of cific carbamoyl B PRODUCTION OF HETASE. AC PO8521). LUCED through a concept the concept and the EMBL.	Profit institutions Institutions as its content is in this statement is not removed. Usage by and for continues alicense agreement (See http://www.isb-sib.ch/arail to license@isb-sib.ch).  AAA33610.1;
n. Sor Sor THE P	are Usag tp:/ Lp:/ 2;
24 AA. date) update) 5'region. cotina; S cotina; S inine-spe inine-spe ATE SYNTH EPTIDE (A	nstitute. There are no ions as long as its case not removed. Usage greement (See http://ww.sb-sib.ch).  Sp-sib.ch).  BFF33F527EAD244C CRC64; Score 28.5; DB 1; Len Pred. No. 2.8e+02;  Mismatches 3; In
Created) Last sequence update) Last annotation update) Arotein in ARG-2 5'region anycota; Pezizomycotina; eae; Neurospora.  Tages, Neurospora.  Tages, Structure and arganine-2 locus. Structure arg-2 locus. Structure seep.  Tages Bannit of arginine-3 argument in special argument argument in a profiture of Bioinformatic	ritute. In the sal long not removed eement (See eement (See isb.ch). The sal long of sal long
STANDARD; PRT; Rel. 19, Created) Rel. 19, Last sequence up Rel. 40, Last annotation assa.  assa. ngi; Ascomycota; Pezizom Sordariaceae; Neurospora 41; IN.A. 10.3 PubMed=2141606; Sachs M.S., Yanofsky C.; ra crassa arg-2 locus. Sit the small subunit of argueraes. ")  1. 265:10981-10987(1990).  AY HAUE A REGULATORY RESPECTEIC CARBAMOYL PHOSPR SPACIFIC CARBAMOYL PHOSPR Y: TO YEAST CPAI LEADER INT. IS COPYLIGHT. IN SWISS INSTITUTE of Bioin SWISS Institute SWISS Institute of Bioin SWISS Institute SWISS In	touce to as to a co to
PRT;  1)  Squency Inotat: In AR( PEZi: Eurospo 1606; 1606; 17(199) 17(199) 17(199) 17(199) 17(199) 17(199) 17(199) 17(199) 17(199)	instructions of the control of the c
D; Created) Last seclast seclast seclast seclast ammyocta; eae; Neueran july yanof arg-2 last subuni; 1 subuni; 3 REGULOSST CRRAMOU	Tantantantantantantantantantantantantanta
ARD; Cre Lase Droot Comy Comy Comy Comy Comy Com	ormatics institute tatement license license 11.00 mm; 27.1%; 53.8%; ative
STANDARD;  1. 19, Cr 1. 19, La 1. 40, La 1. 40, La 8 kDa pro 8a, Romy i, Ascomy i, Ascomy i, Ascomy i, Ascomy in Asc	1 15 IL 122 IL 123 IL 1
STA (Rel. (R	mon-profit insend this statem and this statem and this statem meanle to lice and protein.  24, BAA33610.1  24, BAA33610.1  24, AA, 2779  27. imilarity 53.  7 Conservative FSD-LWKLL 15          :    LSDHLWRAL 22
NEUCR Y3KD	use by non-profit institutes by non-profit institutes are the statement entities requires a license entities requires a license or send an email to license entities facts. J. AAA33610.1;
NEUCR Y3XD NEUCR 01-AUG-1991 0	use by modified would file of the property of
NEUCR STANDARD; PRT; 24 AA.  192702. 101-MUG-1991 (Rel. 19, Created) 101-AUG-1991 (Rel. 19, Last sequence update) 10-OCT-2001 (Rel. 19, Last sequence update) 10-OCT-2001 (Rel. 140, Last annotation update) 10-OCT-2001 (Rel. 140, Last annotation update) 10-OCT-2001 (Rel. 140, Last annotation update) 10-AUG-1991 (Rel. 140, Last annotation update) 10-AUG-1	use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to licenseeisb-sib.ch).  EMBL; J05512, AAA33610.1; - EMBL; J05512, AAA; 2779 MW; 8FF33F527EAD24 Query Match  27.1%; Score 28.5; D Best Local Similarity 53.8%; Pred: No. 2.8e Matches 7; Conservative 2; Mismatches  4 SQETFSD-LWKLL 15  10 SQDYLSDHLWRAL 22
RESULT 1 Y3KD NEUCR 11D Y3KD DAC P2270 DAT 01-AU DT 101-AU DT 101-	Due 3es 1at
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HTF_HELZE
RESULT 4
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                                                                      MEDLINE=93176159; PubMed=8439313;
Menegazzi P., Guzzo F., Baldan B., Mariani P., Treves S.;
"Purification of calreticulin-like protein(s) from spinach leaves.";
Blochem. Blophys. Res. Commun. 190:1130-1135(1993).
                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Caryophyllidae, Caryophyllales, Chenopodiaceae, Spinacia.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Iridaceae;
                                                               STRAIN=CV. GOLDEN HARVEST, and CV. PROF. BLAAUW; TISSUE=Bulb;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24.8%; Score 26; DB 1; Length 20; 44.4%; Pred. No. 5.5e+02; Live 2; Mismatches 3; Indels
                                                                                                                                                                                                                          Score 27; DB 1; Length 20;
Pred. No. 3.9e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- SUBCELLULAR LOCATION: Endoplasmic reticulum lumen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001580; Calreticulin.
PROSITE; PS00803; CALRETICULIN 1; PARTIAL.
PROSITE; PS00804; CALRETICULIN 2; PARTIAL.
PROSITE; PS00805; CALRETICULIN REPEAT; PARTIAL.
Endoplasmic reticulum; Calcium-binding; Glycoprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 AA; 2645 MW; 00FAB4C9DEEDCB0F CRC64;
                                                                                                                                                                                                      20 AA; 2246 MW; DE08DA025FD17D56 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- PTM: GLYCOSYLATED.
-!- SIMILARITY: BELONGS TO THE CALRETICULIN FAMILY.
PIR; PC1241; PC1241.
PIR; PC1240; PC1240.
                                                                                                                                                                                                                                                                                                                                                                   01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                               20 AA.
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                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                           25.7%;
                                                                                                                                                                                                                                                                                                                                                                                                              Spinacia oleracea (Spinach).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 44.4
Matches 4; Conservative
                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                    Calreticulin (Fragment)
                                                                                                                                                                                                                                     Best Local Similarity
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                              NCBI_TaxID=35876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=3562;
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6 PLSEET 11
                                                                                                                                                                                                                                                                      2 PLSQET 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rissum=Leaf;
                                                                                                                                                                                                                                                                                                                                               SPIOL
                                                                                                                                                                                             NON TER
SEQUENCE
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                                                      SEQUENCE
                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                           RESULT 3
CRTC_SPIOL
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Heliothis zea (Corn earworm) (Bollworm).

Eukaryota, Metazoa, Arthropoda, Mandibulata, Pancrustacea, Hexapoda, Insecta, Petrygota, Neopeera, Endopterygota, Lepidoptera, Glossata, Ditrysia, Noctuoidea, Noctuidae, Heliothinae, Helicoverpa.
                                                                                                                                                                  Papio sp. (Baboon).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Cytochrome P450 3A5 (EC 1.14.11) (CYFIIIA5) (P450 FA) (6-beta-testosterone hydroxylase) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                 ..
                                                                                                                                                                                                                                                                                                                                                                                      Score 24; DB 1; Length 10;
Pred. No. 5.3e+02;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                       PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                8E70367865A5B9D1 CRC64;
                     01-AUG-1990 (Rel. 15, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 AA.
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                                                           (HeZ-HRTH).
                                                                                                                                                                                                                                                                                                                                                     AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                        22.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                      62.5%;
                                                                                                                                                                                                                                                                                         PIR; A31571; A31571.
InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation.
MOD RES 10 10
                                                           Hypertrehalosaemic hormone
                                                                                                                                                                                                                                                                                                                                                                                                                 5, Conservative
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STANDARD:
                                                                                                                                                        TISSUE=Corpora cardiaca;
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NCBI_TaxID=61183;
                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 OLTFSSGW 8
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P80056;
 HTF HELZE
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Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Caryophyllidae, Caryophyllales, Chenopodiaceae, Spinacia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           characterization.";
J. Biol. Chem. 273:6710-6716(1998).
-!- SUBCELLULAR LOCATION: Chloroplast; within the thylakoid lumen.
Chloroplast; Thylakoid.
23 23
                      PROSITE; PS00086; CYTOČHROME_P450; PARTIAL.
Oxidoreductase; Monooxygenase; Electron transport; Membrane; Heme;
Microsome; Endoplasmic reticulum.
NON TER 20 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE, AND REVISIONS TO 13 AND 15.
Schubert M., Peterson U., Funk C., Schroeder W.P., Kieselbach T.;
"The chloroplast lumen from Arabidopsis thaliana.";
Submitted (JUL-2001) to the SWISS-PROT data bank.
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01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Benzaldehyde dehydrogenase [NAD+] I (EC 1.2.1.28) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Acinetobatter genomosp. 11.
Bacteria; Proteobacteria; gamma subdivision; Moraxellaceae;
Acinetobacter.
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Pred. No. 1.3e+03;
1; Mismatches 1; Indels
                                                                                                                                                  Score 24; DB 1; Length 20;
Pred. No. 1.1e+03;
                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Andersson B., Schroeder W.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                10-MAY-2000 (Rel. 39, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
17 ylakoid lumenal 17.4 kDa protein (P17.4) (Fragment).
Spinacia oleracea (Spinach).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-98175931; PubMed=9506969;
Kieselbach T., Hagman A., Andersson B., Schroeder W.E
"The thylakoid lumen of chloroplasts - isolation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23 AA; 2571 MW; 022C92717AE9AF2D CRC64;
                                                                                                        SEQUENCE 20 AA; 2226 MW; 3FEF8B6E62BC0F36 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=ATCC 11171 / NCIB 8250 / CIP 63.46 / B94;
MEDLINE=91113163; PubMed=1989592;
                                                                                                                                                                                                                                                                                                                                                                                           23 AA.
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InterPro; IPR001128; Cytochrome P450.
                                                                                                                                                                                                                                                                                                                                                                                              PRT;
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                                                                                                                                                                           Best Local Similarity
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P46365;
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SEQUENCE
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TL17_SPIOL
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"Comparison of benzyl alcohol dehydrogenases and benzaldehyde dehydrogenases from the benzyl alcohol and mandelate pathways in Acinetobacter calcoaceticus and from the TOL-plasmid-encoded toluene pathway in Pseudomonas putida. N-terminal amino acid sequences, amino acid compositions and immunological cross-reactions."; Biochem. J. 273:99-107(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Locusta migratoria (Migratory locust).
Eukaryota, Metazoa, Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Eukaryota; Metazoa, Arthropoda; Mandibulata; Pancrustacea; Caelifera;
Acridomorpha; Acridoidea; Acrididae; Oedipodinae; Locusta.
NCBI_TaxID=7004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Siegert K.J.;
Submitted (DEC-1998) to the SWISS-PROT data bank.
-! FUNCTION: PROBABLY INVOLUED IN THE REGULATION OF LOCUST
INTERMEDIARY METABOLISM, BEHAVIOR AND/OR DEVELOPMENT.
-!- SIMILARITY: SOME SIMILARITY TO THE AKH / HRTH / RPCH FAMILY.
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-!- SIMILARITY: BELONGS TO THE ALDEHYDE DEHYDROGENASES FAMILY.
InterPro; IRR002086; Aldehyde dehydr.
PROSITE; PS00070; ALDEHYDE DEHYDR.CYS; PARTIAL.
PROSITE; PS00687; ALDEHYDE DEHYDR.GLU; PARTIAL.
Aromatic hydrocarbons catabolism; Oxidoreductase; NAD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22.9%; Score 24; DB 1; Length 23; 23.1%; Pred. No. 1.3e+03; ive 3; Mismatches 7; Indels
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Pred. No. 7.6e+02;
0; Mismatches 3; Indels
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01-AUG-1988 (Rel. 08, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
CPAl leader peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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nes 5; Conservative
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1 PNIQTKIIEQIWK 13
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3est_Local Similarity 50.03
Aatches 4; Conservative
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MOD RES 13 13
UNSURE 8 8
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P80181;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
                                                                                                                                                                                                                                                                                                              "Sequence and analysis of a 36.2 kb fragment from the right arm of yeast chromosome XV reveals 19 open reading frames including SNF2 (5'end), CPA1, SLY41, a putative transport ATPase, a putative ribosomal protein and an SNF2 homologue.";
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                                                                                                                                                       MEDLINE=85101411; PubMed=3881260;
Werner M., Feller A., Pierard A.;
"Nucleotide sequence of yeast gene CP Al encoding the small subunit
of arginine-pathway carbamoyl-phosphate synthetase. Homology of the
deduced amino acid sequence to other glutamine amidotransferases.";
Eur. J. Biochem. 146:371-381(1985).
 Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                   Yeast 13:479-482(1997).
-!- SIMILARITY: TO A HYPOTHETICAL PROTEIN IN ARG2 5'REGION FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
                                                  SEQUENCE FROM N.A.
MEDLINE=87215944; PubMed=3555844;
Werner M., Feller A., Messenguy F., Pierard A.;
The leader peptide of yeast gene CPAI is essential for the translational repression of its expression.";
Cell 49:805-813(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21.4%; Score 22.5; DB 1; Length 25; 50.0%; Pred. No. 2.4e+03; tive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                              STRAIN=S288c / FY1679;
BEDLINES-7298310; PubMed=9153758;
Poirey R., Cziepluch C., Tobiasch E., Pujol A., Kordes E.,
Jauniaux J.-C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 AA; 2979 MW; 01D494C8AFEAE4AA CRC64;
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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EMBL; X01764; CAA25904.1; -.
EMBL; Z75210; CAA99620.1; -.
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PIR; S25434; S25434.
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NCBI_TaxID=6523;
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                          NCBI_TaxID=4932;
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TISSUE=Ganglion;
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SEQUENCE 25 A
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P80178;
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Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
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Bukaryota, Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Johnsen A.H., Rehfeld J.F.;
"LymnaDFamides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
MEDLINE=93238777; PubMed=8477756; Johnsen A.H., Rehfeld J.F., "LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaes stagnalis. Clue to cholecystokinin immunoreactivity in
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                                                                                                                      invertebrates?";

Eur. J. Biochem. 213:875-879(1993).

-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.

BIR; 532471; 532471.

Neuropeptide; Amidation.

MOD_RES 13 13 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; S32472; S32472.
                                                                                                                                                                                                                                                                                                                                                          21.0%; Score 22; DB 1; Length 13; 50.0%; Pred. No. 1.4e+03; tive 1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                  9CA07BA3F5D5B455 CRC64;
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MEDLINE=93238777; PubMed=8477756;
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01-JUL-1993 (Rel. 26, Last semi
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                                                                                                                                                                                                                                                                                                  SEQUENCE 13 AA; 1519 MW;
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Coleman M.E., Pan Y.-C.E., Etherton T.D.;
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Bukaryota, Metazoa, Mollusca, Gastropoda, Pulmonata, Basommatophora,
                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Ganglion;
MEDLINE-39238777; PubMed-8477756;
MDDLINE-39238777; PubMed-8477756;
MDDLINE-39238777; PubMed-8477756;
"LymnapFanides, a new family of neuropeptides from the pond snail, Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
            Johnsen A.H., Rehfeld J.F.; "LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Cetartiodactyla, Suina, Suidae, Sus.
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01-MAR-1992 (Rel. 21, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Insulin-like growth factor binding protein 4 (IGFBP-4) (IBP-4)
(IGF-binding protein 4) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eur. J. Biochem, 213:875-879(1993).
-i- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY. PIR; S32475; S12475.
MULTOPEPTIG; Amidation.
MOD RES 13 13 AMIDATION
                                                 invertebrates?";
Eur. J. Biochem. 213:875-879(1993).

Eur. J. Biochem. 213:875-879(1993).

-i - SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.

Neuropeptide; Amidation.

MOD RES. 13 AMIDATION.

SEQÜENCE 13 AA; 1503 MW; 9CA07BBB56D5B455 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 22; DB 1; Length 13;
Pred. No. 1.4e+03;
1; Mismatches 3; Indels
                                                                                                                             Length 13;
                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              9CA07BBB56D5A5A5 CRC64;
                                                                                                                           Score 22; DB 1; L
Pred. No. 1.4e+03;
1; Mismatches 3;
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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       MEDLINE=93238777; PubMed=8477756;
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50.0%;
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50.0%;
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                                                                                                                                                    Conservative
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NCBI_TaxID=6523;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 LSQETFSD 10
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P80182;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Structure, synthesis," and molecular cloning of dermaseptins B, a family of skin peptide antibiotics.";
J. Biol. (Dhem. 273:14680-14697(1998).
-:- FUNCTION: POSSESSES A POTENT ANTIMICROBIAL ACTIVITY AGAINST GRAM-POSITIVE AND GRAM-NEGATIVE BACTERIA. PROBABLY ACTS BY DISTURBING MEMBRANE FUNCTIONS WITH ITS AMPHIPATHIC STRUCTURE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-98278974; PubMed-9614066;
Charpentier S., Amiche M., Mester J., Vouille V., Le Caer J.-P.,
Nicolas P., Delfour A.;
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                                                                                                                                                                                                                                                                                                                                                                                               Length 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -i- SUBCELLULAR LOCATION: Secreted.
-i- SIMILARITY: BELONGS TO THE DERMASEPTIN FAMILY.
Antibictic; Multigene family; Amphibian skin; Amidation.
MOD RES
SEQUENCE 24 AA; 2665 MW; E5987D7F50E08F4F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Indels
                                                                                                                                                                                                                                                                                                                                                       SEQUENCE 16 AA; 1799 MW; 40988840096655E2 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                             21.0%; Score 22; DB 1; I
66.7%; Pred. No. 1.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-Ermaseptin BVI (Dermaseptin B6)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Phyllomedusa bicolor (Two-colored leaf frog).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Mismatches
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Best Local Similarity 50.00
                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Conservative
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

February 12, 2003, 11:12:22 ; Search time 20.5833 Seconds (without alignments) 88.739 Million cell updates/sec

US-09-403-440A-2

105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

**BLOSUM62** Scoring table:

Gapop 10.0 , Gapext 0.5

283224 seqs, 96134422 residues Searched:

4984 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 73:\*
1: pir1:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

genome polyprotein c-myc protein - ch protein phosphatas hypothetical prote gamma-crystallin I cytochrome P450 3A antifungal 2S stor alpha-1-antitrypsi Ig heavy chain DJ hypothetical prote Tha p 1 - Thaumeto Ig heavy chain DJ hypertrehalosemic/ chaperonin 10 homo base nonspecific a adipokinetic hormo calcium-binding pr translation elonga pollen allergen Po testosterone 6beta insulin-like growt benzaldehyde dehyd phosphoprotein pho lectin chain B - I H+-exporting ATPas ribulose-bisphosph (cytosine-5-)water-soluble 35K Description SUMMARIES T42442 B42224 D25865 PC1240 A61506 PH1334 A59396 A60728 A53592 A34921 PH1326 A31571 D42842 A35105 PA0110 A60372 PC2347 S35641 S11545 A24244 S41747 S21176 S13298 T42259 DB Query Match Length 30 28.5 Score Result 

3

Gaps

9

1 PPLSQETFSDLWK---LLP 16 

RESULT 3

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T42442

protein kinase (EC	cytochrome-c oxida	phosphorotein pho	Gene CDA1 leader n	endo-1.4-beta-xvla	hypothetical prote	lymnaDFamide 1 - q	lymnaDFamide 2 - g	lymnaDFamide 4 - g	lymnaDFamide 5 - g	L-2.4-diaminobutor	insulin-like growt	hydroxyorol ine-ric	nlasma proteinase	intracryotalline o	nucleoside-diphosp
S20410	S65379	T42257	\$25434	870615	S16324	S32471	S32472	S32474	832475	B44854	JH0517	S59481	A60812	A56894	868023
7	7	7	~	~	7	N	~	~	~	~	7	~	N	~	7
15	23	24	25	2	σ	13	13	13	13	14	16	17	20	20	21
21.9	21.9	21.9	21.4	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0
~	23	23	22.5	22	22	22	22	22	22	22	22	22	22	22	22
5															

#### ALIGNMENTS

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genome polyprotein (clone L3/S2) - Skalica virus (fragment)
C;Species: Skalica virus
C;Species: 9-Mar-1997 #sequence_revision 19-Dec-1997 #text_change 17-Nov-2000
C;Accession: B61409
R;Guirakhoo, F.; Heinz, F.X.; Mandl, C.W.; Holzmann, H.; Kunz, C.; Gresikova, M. J. Gen. Virol. 72, 333-338, 1991
A;Title: The relationship between the flaviviruses Skalica and Langat as revealed by mon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Westaway, D.; Payne, G.; Varmus, H.E.
Proc. Natl. Acad. Sci. U.S.A. 81, 843-847, 1984
A;Title: Proviral deletions and oncogene base-substitutions in insertionally mutagenized
A;Reference number: I50387; MUID:84144799; PMID:6322173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  c-myc protein - chicken (fragment)
C;Species: Gallus gallus (chicken)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 21-Jul-2000
C;Accession: IS0387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-24 <WES>
A;Cross-references: GB:K01440; NID:g212357; PIDN:AAA48964.1; PID:g212358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
                                                                                                                                                                                                                                             A Reference number: A61409, MUID:91132129; PMID:1847173
A;Accession: B61409
A;Status: preliminary; not compared with conceptual translation
A;Molecule type: genomic RNA
A;Residues: 1-19 <GUI>
C;Superfamily: yellow fever virus genome polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      32.4%; Score 34; DB 2; Length 19; ilarity 41.7%; Pred. No. 81; Conservative 3; Mismatches 4; Indels
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2 PPVTYRTGTDCW 13
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les 5; Conserv
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B61409
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C;Accession: A60728
R;Pineau, T.; Galtier, P.; Bonfils, C.; Derancourt, J.; Maurel, P.
Bicchem. Pharmacol. 39, 901-909, 1990
A;Title: Purification of a sheep liver cytochrome P-450 from the P450IIIA gene subfamily A;Reference number: A60728; MUID:90179800; PMID:2310415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         datifugal 2S storage albumin large chain - radish (fragment)
C;Species: Raphanus sativus (radish)
C;Species: Raphanus sativus (radish)
C;Date: 27-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 12-Apr-1995
C;Accession: D42842
R;Terras, F.R.; Schoofs, H.M.; De Bolle, M.F.; Van Leuven, F.; Rees, S.B.; Vanierleyden, J. Blol. Chem. 267, 15301-15309, 1992
A;Title: Analysis of two novel classes of plant antifungal proteins from radish (Raphanu A;Reference number: A42842; MUID:92348373; PMID:1633777
                                                                                                                                                                                                                                submitted to the Protein Sequence Database, September 2001
A;Description: Isolation and characterization of a major allergen of the pine Frocession
A;Reference number: A59396
A;Accession: A59396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A,Gene: CYP3A
C;Superfamily: human cytochrome P450 CYP3A5; cytochrome P450 homology
C;Keywords: electron transfer; endoplasmic reticulum; heme; monooxygenase; oxidoreductas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: protein
A;Residues: 1-20 <PIN>
C;Comment: This cytochrome P450 isozyme is a member of the P450IIIA family but is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytochrome P450 3A, troleandomycin-induced - sheep (fragment)
N;Contains: oxidoreductase (EC 1.-.-.)
C;Species: Ovis sp. (sheep)
C;Date: 14-May-1993 #sequence_revision 14-May-1993 #text_change 05-Mar-1999
                                                                                                                                       C;Spēcies: Thaumetopoea pityocampa
C;Date: 03-Jun-2002 #sequence_revision 03-Jun-2002 #text_change 03-Jun-2002
C;Accession: A59396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24.8%; Score 26; DB 2; Length 18; 38.5%; Pred. No. 1.2e+03; tive 3; Mismatches 5; Indels
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46.7%; Pred. No. 1.4e+03;
tive 1; Mismatches 3; Indels
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A;Note: sequence extracted from NCBI backbone (NCBIP:109925)
                                                                                                            Tha p 1 - Thaumetopoea pityocampa (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A,Experimental source: L5
A,Note: IgE-binding protein, major allergen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 46.7
Matches 7; Conservative
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Best Local Similarity 38.5'
Matches 5; Conservative
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A;Molecule type: protein
A;Residues: 1-20 <TER>
                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: protein A; Residues: 1-18 < MON>
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                                                        RESULT 6
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                                                                                     C,Accession: T4242
R;Zeke, T.; Gergely, P.; Dombradi, V.
Submitted to the EMBL Data Library, July 1996
A;Description: The catalytic subunits of Ser/Thr protein phosphatases from Caenorhabditi
A;Reference number: Z22025
A;Accession: T42442
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C,Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 21-Jul-2000 C;Accession: B42224  
R;Orbach, M.J.; Sachs, M.S.; Yanofsky, C. J. Biol. Chem. 265, 10981-10987, 1990 A;Title: The Neurospora crassa arg-2 locus. Structure and expression of the gene encodin A;Reference number: A42224; MUID:90293033; PMID:2141606
                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: EMBL:Z77728; PIDN:CAB01287.1
C;Superfamily: phosphoprotein phosphatase; phosphoesterase core homology; phosphoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C,Accession: D25865
R,Chiou, S.H.; Chen, S.W.; Lo, T.B.
R;Chiou, S.H.; Chen, S.W.; Lo, T.B.
R;EBS Lett. 209, 107-110, 1986
A;Title: The amino-terminal sequences of four major carp gamma-crystallin polypeptides
A;Reference number: A91369; MUID:87105907; PMID:3803569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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A;Residues: 1-23 <CHI>
A;Note: the residues 3-11e, 15-Asn, 17-Asp, 18-Cys, and 20-His were also found
C;Superfamily: beta-crystallin
C;Reywords: duplication
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypothetical protein (arg-2 5' region) - Neurospora crassa
C;Species: Neurospora crassa
C;Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gamma-crystallin IV - common carp (fragment)
C;Species: Cyprinus carpio (common carp)
C;Date: 02-Jun-1988 #sequence_revision 02-Jun-1988 #text_change 21-Nov-1997
                       C:Species: Caenorhabditis elegans
C:Species: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 02-Jun-2000
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A;Residues: 1-24 <ORB>
A;Cross-references: GB:J05512; NID:g168864; PIDN:AAA33610.1; PID:g168866
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 28.6%; Score 30; DB 2; Length 24; Best Local Similarity 58.3%; Pred. No. 4.2e+02; Matches 7; Conservative 1; Mismatches 4; Indels
protein phosphatase - Caenorhabditis elegans (fragment)
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Best Local Similarity 53.0.
The conservative
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Best Local Similarity 75.09
Matches, 6; Conservative
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A; Residues: 1-24 <ZEK>
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R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1334
A;Molecule type: DNA
A;Residues: 1-23 «WAS>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nypothetical protein - Neurospora crassa mitochondrion (fragment)
C;Species: mitochondrion Neurospora crassa
C;Species: mitochondrion Neurospora crassa
C;Species: mitochondrion Neurospora crassa
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 07-Dec-1999
C;Accession: A35105
R;Saville, B.J.; Collins, R.A.
Cell 61, 685-696, 1990
A;Title: A site-specific self-cleavage reaction performed by a novel RNA in neurospora A;Reference number: A35105; MUD: 90263093; PMID: 2160856
A;Accession: A35105
A;Accession: A35105
A;Accession: A35105
A;Accession: A35105
C;Genetics:
C;Genetics:
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pollen allergen Poa-pI - Kentucky bluegrass (fragment)
C;Species: Poa pratensis (Kentucky bluegrass)
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Best Local Similarity 33.33
Watches 4; Conservative
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Best Local Similarity 54.59
Laga 6; Conservative
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A, Genetic code: SGC3
C, Keywords: mitochondrion
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Comp. Biochem. Physiol. B 100, 293-296, 1991
A;Title: Purification, characterization, and acute phase response of plasma alpha-1-anti
A;Reference number: A61506; MUID:92191572; PMID:1724745
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Cispecies: Spinacia oleracea (spinach)
Cispecies: Spinacia oleracea (spinach)
Cispecies: 30-sep-1993 #sequence_revision 30-sep-1993 #text_change 23-Aug-1997
Cist Cispecia in Prof. 2012 Cist i
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C;Species: Mesocricetus auratus (golden hamster)
C;Date: 15-Oct.1994 #sequence_revision 15-Oct.1994 #text_change 11-May-2000
C;Accession: A61506
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1334
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                                                                     Query Match 24.8%; Score 26; DB 2; Length 20; Best Local Similarity 45.5%; Pred. No. 1.4e+03; Matches 5; Conservative 3; Mismatches 3; Indels
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C; Superfamily: wheat alpha-amylase inhibitor
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C,Superfamily: calreticulin
C,Keywords: calcium binding; metal binding
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Matches 6; Conservative
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A,Residues: 1-20 <ME2>
A,Experimental source: leaf
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-20 <AME>
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5 EERFEDGWE 13
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H-exporting ATPase (EC 3.6.3.6) 24k chain - Thermus aquaticus (fragment)
C;Species: Thermus aquaticus
C;Species: Thermus aquaticus
C;Date: 01-Dec-1995 #sequence_revision 01-Dec-1995 #text_change 03-Jun-2002
R;Vokcyama, K.; Akbane, Y.; Ishii, N.; Yoshida, M.
J. Biol. Chem. 269, 12248-12253, 1994
J. Biol. Chem. 269, 12248-12253, 1994
J. Biol. Chem. 269, 12248-12253, 1994
J. Reference number: A53592; MUID:94216345; PMID:8163530
A;Accession: A53592
A;Accessi
C;Date: 11-Feb-1993 #sequence_revision 11-Feb-1993 #text_change 09-Jun-2000
C;Accession: A60372; A33086
R;Ekramoddoullah, A.K.M.
Int. Arch. Allergy Appl. Immunol. 93, 371-377, 1990
A;Title: Two-dimensional gel electrophoretic analyses of Kentucky bluegrass and rye grasence of Poa p I allergen.
A;Reference number: A60372; MUID:91317571; PMID:2101126
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C;Keywords: hydroxyproline; pollen
F;5,8/Modified site: hydroxyproline (Pro) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

23.8%; Score 25; DB 2; Length 20;
Best Local Similarity 36.4%; Pred. No. 2e+03;
Matches 4; Conservative 1; Mismatches 6; Indels
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23.8%; Score 25; DB 2; Length 20;
Best Local Similarity 46.2%; Pred. No. 2e+03;
Matches 6; Conservative 3; Mismatches 4; Indels
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A;Molecule type: protein
A;Residues: 1-20 <EKR>
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February 12, 2003, 10:43:13; Search time 30 Seconds (without alignments) 18.635 Million cell updates/sec
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/pcTUS COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/pcTUS COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                              262574 seqs, 29422922 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                            OM protein - protein search, using sw model
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1 PPLSQETFSDLWKLLPENG 19
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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	Sequence 13, Appl Sequence 1, Appli Sequence 1, Appli Sequence 20, Appli Sequence 20, Appli Sequence 21, Appli Sequence 21, Appli Sequence 21, Appli Sequence 21, Appli Sequence 6, Appli Sequence 6, Appli Sequence 6, Appli Sequence 17, Appli Sequence 17, Appli Sequence 17, Appli Sequence 46, Appli Sequence 17, Appli Sequence 46, Appli Sequence 46, Appli Sequence 17, Appli Sequence 17, Appli	15,
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ength DB	4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	11 1
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Sequence Seq	o, AND	0;
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S-08 424-957-23 S-08 424-957-28 S-09 424-957-28 S-09 424-957-40 S-09 035-686-28 S-09 035-686-40 S-08 424-957-11 S-08 424-957-11 S-08 424-957-36 S-08 424-957-34 S-08 424-957-34 S-08 424-957-35 S-08 424-957-35 S-08 424-957-35 S-08 424-957-35 S-08 424-957-35 S-08 424-957-36 S-08 424-957-36 S-08 424-957-35 S-08 424-957-35 S-08 424-957-36 S-09 635-686-33	NTS AND On 2.	re 90; DB 4; L, d. No. 4.2e-08; Mismatches 1;
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	cation US/0908197  cation US/0908197  ilin, William  tt, Christine  TION: METHODS OF  TION: METHODS OF  TION: METHODS OF  TION: METHODS  METHODS OF  TION: MATHODIES  METHODS OF  TION: MATHODIES  MODERS:  LE FORM:  Disker of  TERN:  UMBER: 0S/09/081  12-MAY-1998  N: 435  ON DATA:  UMBER: 0S/09/081  TERN:  UMBER: 06/046,20  12-MAY-1998  N: 435  ON DATA:  UMBER: 06/046,20  12-MAY-1998  N: 435  ON DATA:  UMBER: 30,628  KET NUMBER: 4740  ON INFORMATION:  13-345-6054  -345-1300  EQ ID NO: 13:  TERISTICS:  mind acids	85.7%; 88.9%; ative
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RESULT 4
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Patent No. 5702908
GENERAL INFORMATION
GENERAL INFORMATION
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
RADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREYT: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                Sequence 1, Application US/08277660A
Patent No. 5702908
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 73.3%; Score 77; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Flour Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California STATE: California ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRENT AFFLICATION DATE:

CLASSIPLATION NUMBER: US/08/277,660A
FILING DATE: 20-UL-1994
CLASSIPLATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 781-1989
TELERAX: (415) 781-1989
TELERAX: 910 277299
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
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                          1 PPLSQETFSELWNLLPEN 18
1 PPLSQETFSDLWKLLPEN 18
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US-08-277-660A-1
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COUNTRY:
ZIP: 9411
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US-08-277-660A-1
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Sequence 1, Application US/08424957
| Patent NO. 5770377
| Patent NO. 5770377
| CENBEAL INFORMATION:
| APPLICANT: Land, David P. TITLE OF INVENTION: Interruption of Binding of MDM2 and PS3 TITLE OF INVENTION: Protein and Therapeutic Application Thereof; COMESPONDENCE ADDRESS:
| COMESPONDENCE ADDRESS:
| ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 73.3%; Score 77; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESSEE: Plehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco STRTE: California COUNTRY: United States ZIP: 94111-487 COMPUTER READBLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957 FILING DATE: 19-APR-1995 CLASSIFICATION: PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660 FILING DATE: 20-UU-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                  SCHWARE SILE. 1. 2007/130 SCHWARE BEACHIN Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A FILING DATE: 20-UUL-1994
CLASSIFICATION: 514
CLASSIFICATION: 514
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A-61228/WHD
                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: A-60244/WHD TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMEDUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 amino acids TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE, DOCKET NUMBER: A-6
TELECOMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPAN: (415) 781-1989
TELEFAX: (415) 398-3249
TELEEX: 910 27729
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKL 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-277-660A-4
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GENERAL INFORMATION:
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flahr, Hobbach, Test, Albritton & Herbert

STREET: San Francisco

CITTLE California

STATE: California

COUNTRY: United States

CONTRY: DATE: CALIFORNIA

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                                                                                                                                                                                                                                       73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
PRIOR DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: 3,61228/WHD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 1, Application US/09035686
; Patent No. 6153391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15 amino acids
            15 amino acids
                                                                                                                                                                                                                                                                                                                        14; Conservative
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Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     5 QETPSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 PPLSQETFSDLWKL 14
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                                                                                                                         TOPOLOGY: unknown

; TOPOLOGY: unknown
US-08-424-957-20
                                                                                                                                                                                                                                                                               Best Local Similarity
                                            TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 5
US-08-424-957-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
TELEX: 9
                                                                                                                                         JUS-08-424-957-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-035-686-1
LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                                        Matches
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Gaps
APPLICANT: Ficksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
OCRRESPONDENCE ADDRESS:
ADDRESSE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   73.3%; Score 77; DB 4; Length 15; 100.0%; Pred. No. 3.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CURRANT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
FILING DATE: 19-APR-1995
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 19-APR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Dreeger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/POCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ....
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TELEEX: 910 277-299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 OETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: unknown
US-09-035-686-1
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US-09-035-686-20
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STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                73.3%; Score 77; DB 4; Length 15; 100.0%; Pred. No. 3.8e-06; tive 0; Mismatches 0; Indels
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-AR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 781-1989
TELEFAX: (415) 781-1989
TELEFAX: 910 277299
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: amino acide
TYPE: ATTORNEY: STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A-60244/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: A-6
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.0
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       : 15 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 PPLSQETFSDLWKL 15
                                                                                                                                                                                                                                                                                                                                                      unknown
                                                              CLASSIFICATION:
                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
US-09-035-686-20
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Patent No. 5770377

GENERAL INFORMATION:

APPLICANT: David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert

STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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                                                            70.5%; Score 74; DB 1; Length 15; 93.3%; Pred. No. 1.2e-05; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 70.5%; Score 74; DB 1; Length 15; 93.3%; Pred. No. 1.2e-05; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: A-61228/WHD TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE PORM:
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-09-035-686-21
; Sequence 21, Application US/09035686
; Patent No. 6153391
GENERAL INFORMATION:
APPLICANT: Pickeley, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ 1D NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415) .--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                               Query Match
Best Local Similarity 93.33
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PLSOETFSGLWKLLP 15
                                                                                                                                                           2 PLSQETFSDLWKLLP 16
                                                                                                                                                                                        1 PLSQETFSGLWKLLP 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: unknown
US-08-424-957-21
; TOPOLOGY:
US-08-277-660A-5
                                                                                                                                                                                                                                                                             RESULT 9
US-08-424-957-21
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Patent No. 6451979
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
MILIAM APPLICANT: Gost, Christine
APPLICANT: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SECURIORES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: Californias
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
0
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                                                                                                                                                                                                                                    ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERALING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
TELERAX: 910 277299
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PLSQETFSGLWKLLP 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 PLSQETPSDLWKLLP 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: unknown
US-09-035-686-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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US-09-081-975-23
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STATE: M
COUNTRY:
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CURRENT APPLICATION DATA:

JOURNAL APPLICATION DATA:

PRILING DATE:

TILING DATE:

TIL
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TELEFAX:
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                                                                                      0; Gaps
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0
                                                                                                                                                                                                                                                                                                                                         APPLICANT: GOUELI, SAID A.
TITLE OF INVENTION: QUANTITATION OF INDIVIDUAL PROTEIN
KINASE ACTIVITY
                                        61.9%; Score 65; DB 3; Length 15; 84.6%; Pred. No. 0.00031; Live 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Ross & Stevens, S.C.
STREET: One South Pinckney Street, P.O. Box 2599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61.9%; Score 65; DB 4; Length 15; 84.6%; Pred. No. 0.00031; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CT-USS5-02856-6
Sequence 6, Application PC/TUS9502856
GENERAL INFORMATION:
APPLICANT: Goueli, Said A
TITLE OF INVENTION: QUANTITATION OF INDIVIDUAL PROTEIN
TITLE OF INVENTION: KINASE ACTIVITY
NUMBER, OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/208,573B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 10-DEC-1998
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: SARA, CHARLES S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 34506.006
TELECOMUNICATION INFORMATION:
TELEPHONE: (608) 257-5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear

MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 6:
US-08-208-573B-6
                                                                                                                                                                                                                                                                            Sequence 6, Application US/08208573B Patent No. 6348310 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (608) 257-9175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 10
                                    Query Match
Best Local Similarity 84.6'
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Wisconsin COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 84.6
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 53701-2599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Madison
                                                                                                                             1 PPLSQETFSDLWK 13
                                                                                                                                                      2 PPLSQEAFADLWK 14
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US-08-208-573B-6
US-09-280-047-6
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APPLICANT: Kaelin, William
APPLICANT: Cost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61.9%; Score 65; DB 5; Length 15; 84.6%; Pred. No. 0.00031; tive 1; Mismatches 1; Indels
COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IEM Compatible
COMPUTER: IEM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: REALESQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 34506.006
TELECOMUNICATION INFORMATION:
TELEPHONE: 608-831-2100
TELEFAX: 608-831-2106
                                                                                                                                                                                  APPLICATION NUMBER: PCT/US95/02856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 22, Application US/09081975; Patent No. 6451979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                       CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 474
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                    9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 608-831-2106
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 84.6
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PPLSQEAFADLWK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
GY: linear
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                                                                                                                                                                                                       FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT-US95-02856-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-081-975-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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; TELER:
; INPORMATION FOR SEQ ID NO: 22:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 18 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-081-975-22

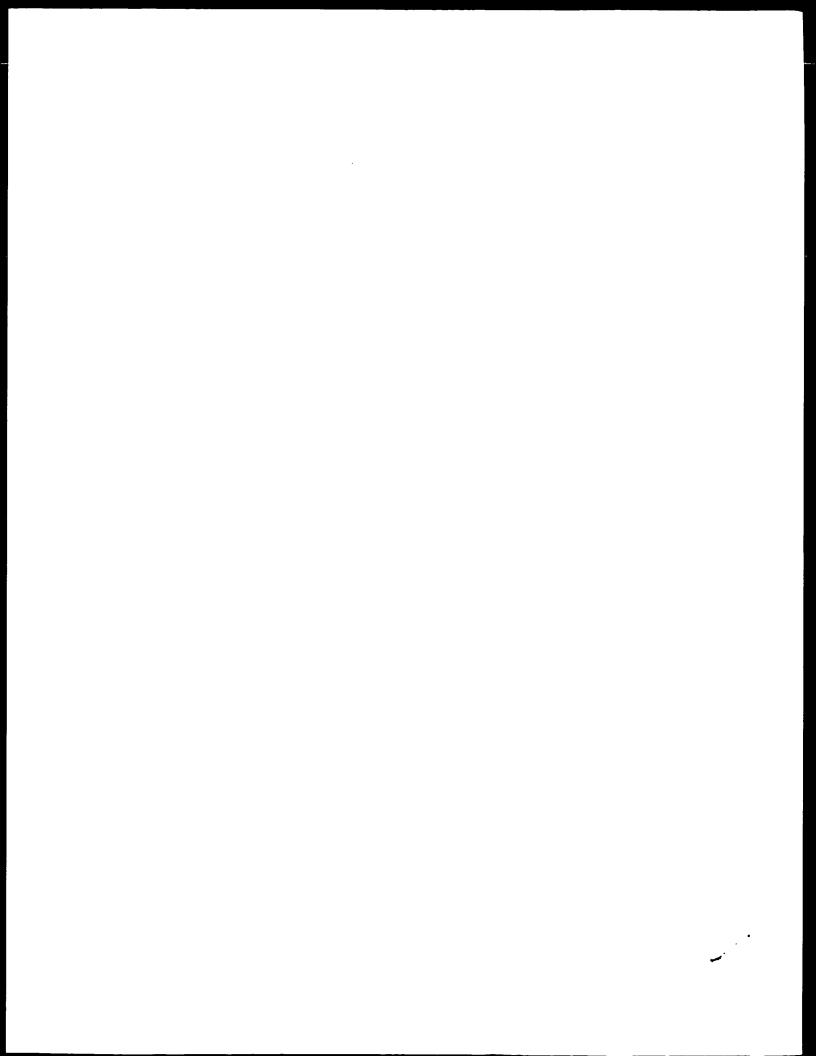
Query Match
Best Local Similarity 76.9%; Pred. No. 0.0024;
Matches 10; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 2 PLSQETFSDLWKL 14

Cy 2 PLSQEFFSDLWKM 15

Db 3 PLSQESFFDLWKM 15

Search completed: February 12, 2003, 10:46:27
Job time: 30 secs
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February 12, 2003, 10:44:28; Search time 56 Seconds (without alignments) 28.428 Million cell updates/sec
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                           OM protein - protein search, using sw model
                                                                                                                                                    Run on:
```

105 1 PPLSQETFSDLWKLLPENG 19 US-09-403-440A-2 Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

425801 seqs, 83787487 residues Searched:

152190 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Pending Patents AA New:\*

1: /cgn2\_6/ptodata/1/paa/USO6\_NEW\_COMB.pep:\*

2: /cgn2\_6/ptodata/1/paa/USO6\_NEW\_COMB.pep:\*

3: /cgn2\_6/ptodata/1/paa/USO8\_NEW\_COMB.pep:\*

3: /cgn2\_6/ptodata/1/paa/USO8\_NEW\_COMB.pep:\*

5: /cgn2\_6/ptodata/1/paa/USO8\_NEW\_COMB.pep:\*

6: /cgn2\_6/ptodata/1/paa/USO8\_NEW\_COMB.pep:\*

7: /cgn2\_6/ptodata/1/paa/USO8\_NEW\_COMB.pep:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

		de			SUMMAKIES	
Result		Query				
No.	Score	Match	Match Length DB	DB	ID	Description
	92	87.6	19	7	US-60-438-805-9	Semience 9 specific
7	81	77.1	15	9	US-10-211-088-141	. 4
m	70	66.7	13	ß	US-09-701-080C-22	
4	70	66.7	13	ß	US-09-701-080C-27	Sequence 27. April
S	59	56.2	11	S	US-09-189-702A-286	286
9	41	39.0	14	9	US-10-049-428-6	9
7	40	38.1	7	Ŋ	US-09-701-080C-12	
œ	35	33.3	14	9	US-10-049-428-7	, ,
σ	31.5	30.0	14	9	US-10-268-332-53	
10	31.5	30.0	16	9	US-10-225-567A-1177	Sequence 1177. An
11	31	29.5	o,	Ŋ	US-09-189-702A-7	
12	31	29.5	12	Ŋ	US-09-281-717A-31	~
13	31	29.5	12	Ŋ	US-09-281-717A-33	3 1
14	31	29.5	12	ហ	US-09-281-717A-35	Sequence 35, Appl
15	31	29.5	12	ß	US-09-830-693B-6	Ġ
16	31	29.5	12	Ŋ	US-09-830-693B-8	à
17	30	28.6	16	9	US-10-348-131-49	Sequence 49. Appl
18	29	27.6	15	9	US-10-169-026-3	<u>س</u>
19	29		16	9	US-10-225-567A-942	σ
20	29	27.6	17	н	PCT-US02-28371-70	70.
21	29	27.6	17	ø	US-10-236-878-70	Sequence 70. Appl
22	29	27.6	18	ß	US-09-552-802B-30	30
23	28.5	27.1	18	S	US-09-632-036E-11	11,
24	28	26.7	7	9	US-10-174-717A-36	36
25	28	26.7	16	7	US-60-427-045-297	297
26	28	26.7	18	9	US-10-283-599-219	219,

TYPE: PRT

	Sequence 25791, A
	US-09-641-528A-25791
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8,7,7,7,7,7,7,7,7,7,8	7
7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	<b>1</b> *

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Gaps
                                                                                                                                                                                                           ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: Peptide
US-60-438-805-9
                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                 Query Match

87.6%; Score 92; DB 7; Length 19;
Best Local Similarity 100.0%; Pred. No. 4.5e-08;
Matches 17; Conservative 0; Mismatches 0; Indels
          Sequence 9, Application US/60438805;
GRERAL INFORMATION:
APPLICANT: KODADEK, THOMAS;
ITILE OF INVENTION: MIXED BLEMENT CAPTURE AGENTS;
FILE REFERENCE: UTSD:935USP1;
CURRENT APPLICATION NUMBER: US/60/438,805;
CURRENT FILING DATE: 2003-01-09;
NUMBER OF SEQ ID NOS: 12;
SEQ ID NO 9;
LENGTH: 19
                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                        2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                        1 PLSQETFSDLWKLLPEN 17
US-60-438-805-9
                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                             RESULT 2
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0; Gaps
Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 56.2%; Score 59; DB 5; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0046; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                         APPLICANT: Sidey, John
APPLICANT: Sidey, John
APPLICANT: Southwood, Scott
APPLICANT: Southwood, Scott
APPLICANT: Boundare, Inc.
TITLE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REPERBENCE: 3963-20019.20
CURRENT APPLICATION NUMBER: US 09/205,713
PRIOR PLING DATE: 1998-11-10
PRIOR PLING DATE: 1998-11-10
PRIOR PLING DATE: 1998-304
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FRASEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Xu, Weiming
APPLICANT: Xu, Weiming
APPLICANT: Liu, Lizhi
TITLE OF INVENTION: Unducible Screen for Drug Discovery
FILE REFERENCE: HO-P02380USO
CURRENT APPLICATION NUMBER: US/10/049,428
CURRENT APPLICATION NUMBER: GB 9918077
PRIOR APPLICATION NUMBER: GB 9918077
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
Indels
.;
0
  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: p53.14 peptide 34.0324
                                                                                                                                                                                                              ; Sequence 286, Application US/09189702A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/10049428; GENERAL INFORMATION: APPLICANT: Charles, Ian G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                    APPLICANT: Sette, Alessandro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 69.2
Matches 9; Conservative
Matches 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWK 13
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                                                 4 SQETFSDLWKLLP 16
                                                                                                   1 SQETFSDLWKLLP 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 LSQETFSDLWK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 LSQETFSDLWK 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: HUMAN
                                                                                                                                                                         RESULT 5
US-09-189-702A-286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-189-702A-286
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US-10-049-428-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 286
LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-049-428-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 6
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1. Sequence 22, Application US/09701080C

1. Sequence 22, Application US/09701080C

2. GENERAL INFORMATION:
CENTRAL INFORMATION:
APPLICANT: UNISTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REPREMENCE: N7347C GW
FILE REPREMENCE: N7347C GW
FILE REPREMENCE: N7347C GW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT RILING DATE: 2001-02-27
PRIOR PLING DATE: 1998-05-26
PRIOR FILING DATE: 1998-05-26
PRIOR PLING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN VET. 2.1
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TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REFERENCE: N3477C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 1998-05-26
PRIOR PILING DATE: 1998-05-26
PRIOR PLILING DATE: 1999-01-05
NUMBER: OF SEQ ID NOS: 36
SOFTWARE: PATCHING Ver: 2.1
SEQ ID NO 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence:derived from p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
                                                                                                                                                                              0; Gaps
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                                                                                                                          Query Match

77.1%; Score 81; DB 6; Length 15;
Best Local Similarity 100.0%; Pred. No. 2e-06;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y Match 66.7%; Score 70; DB 5; Length 13; Local Similarity 100.0%; Pred. No. 9.6e-05; hes 13; Conservative 0; Mismatches 0; Indels
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                                                      CTHER INFORMATION: Binding domain US-10-211-088-141
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          ORGANISM: Artificial sequence
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Sequence 1177, Application US/1022567A

GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Christine C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
FILE REFERENCE: 1920-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOSTWARE: PARENTIN VERSION 3.1
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FILE REFERENCE: D0042A CIP
CURRENT APPLICATION: INMUNE: US/10/268,332
CURRENT PELING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US. 60/235,713
PRIOR FLING DATE: 2000-09-27
PRIOR FLING DATE: 2000-09-27
PRIOR FLING DATE: 2000-09-27
PRIOR FLING DATE: 2001-01-16
PRIOR FLING DATE: 2001-01-16
PRIOR FLING DATE: 2001-01-17
PRIOR FLING DATE: 2001-01-17
PRIOR FLING DATE: 2001-01-18
PRIOR PRIOR APPLICATION NUMBER: U.S. 60/313,171
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-09-26
NUMBER OF SEQ ID NOS: 66
SOFTWARE: PATENTIN VERSION 3.1
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53.3%; Pred. No. 1.6e+02;
tive 2; Mismatches 2; Indels
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GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidnsy, John
APPLICANT: Southwood, Scott
APPLICANT: Bpimmune, Inc.
APPLICANT: Epimmune, Inc.
ITILE OF INTENTION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019.20
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Best Local Similarity 53.3.
Best Local 8; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Matches 7; Conserv
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APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY3, EXPRESSED HIGHI
                                     Sequence 12, Application US/09701080C
GRNERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
FILE REPREMENCE: N/34/7C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 2001-02-27
FRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR FILING DATE: 1998-05-26
PRIOR PLING DATE: 1998-05-26
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SEQ ID NOS: 36
SEQ ID NO 12
LEAGURE 10 10 10 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Charles, Ian G.
APPLICANT: Charles, Ian G.
APPLICANT: Xu, Weiming
APPLICANT: Liu, Lizhi
TITLE OF INVENTION: Unducible Screen for Drug Discovery;
FILE REFERENCE: HO-P02380US0
CURRENT FILING DATE: 2000-07-28
PRIOR APPLICATION NUMBER: US/10/049,428
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          38.1%; Score 40; DB 5; Length 7; 100.0%; Pred. No. 3.5e+05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 7; Conservative
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Matches 7; Conservative
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1 FSDLWKL 7
                      US-09-701-080C-12
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LENGTH: 14
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  RESULT 7
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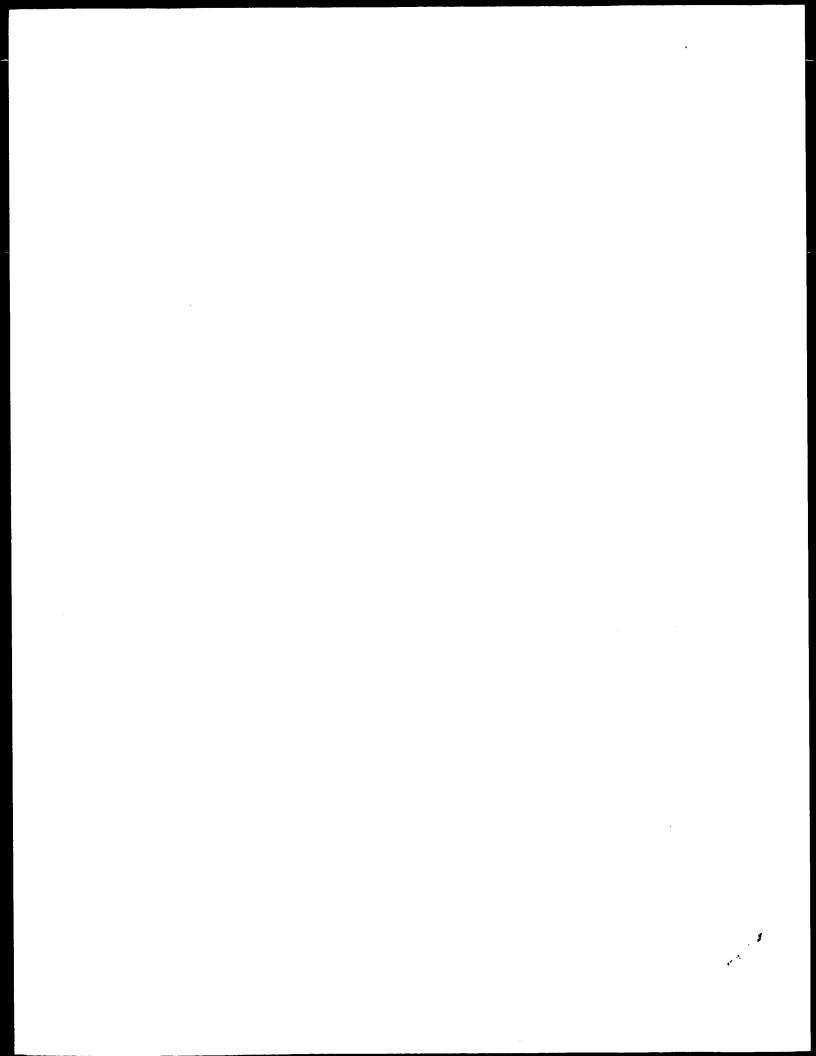
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Sequence 33, Application US/09281717A GENERAL INFORMATION:
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Best Local Similarity 60.v.
Local 6, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: Homo sapiens US-09-281-717A-33
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US-09-281-717A-35
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US-09-830-693B-6
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LENGTH: 12
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APPLICANT: Part weijun
APPLICANT: Fletterick, Robert
APPLICANT: Fletterick, Robert
APPLICANT: Fletterick, Robert
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Robert
APPLICANT: Wagner, Robert
APPLICANT: Wagner, Robert
APPLICANT: Wagner, Robert
APPLICANT: Wagner, Weith
TITLE OF INVENTION: COACTIVACR BINDING
FILE REFERENCE: 9811-008-999
CURRENT APPLICATION NUMBER: US 60/079,956
PRIOR PRILITY DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
SSEQ ID NO 31
LENGTH: 12
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                                                                                                                                                                                                                                                                                              Query Match 29.5%; Score 31; DB 5; Length 9; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 6; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                     ) OTHER INFORMATION: p53.24V9 peptide 1323.02
US-09-189-702A-7
  CURRENT APPLICATION NUMBER: US/09/189,702A CURRENT FILING DATE: 1998-11-10 PRIOR APPLICATION NUMBER: US 08/205,713 PRIOR FILING DATE: 1994-03-04 NUMBER OF SEQ ID NOS: 380 SGYTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (5)..(5)
OTHER INFORMATION: Leu --> Arg (L454R)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (7)..(7)
OTHER INFORMATION: Leu --> Arg (L456R)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; NAME/KEY: MUTAGEN
; LOCATION: (8) .. (8)
; OTHER INFORMATION: Glu -->Lys (E457K)
US-09-281-717A-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 31, Application US/09281717A; GENERAL INFORMATION:
                                                                                                                                                                             TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 60.00
المالية 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Baxter, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PPLSOETFSD 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: MUTAGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: MUTAGEN
                                                                                                                                                                                                                                                                                                                                                                                     13 KLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                             1 KLLPEN 6
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US-09-281-717Ax33
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US-09-281-717A-31
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APPLICANT: Baxter, John
APPLICANT: Baxter, John
APPLICANT: Baxter, John
APPLICANT: Parimont, Beatrice
APPLICANT: Fletterick, Robert
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: COACTIVAOR BINDING
TITLE OF INVENTION: COACTIVAOR BINDING
FILE REFERENCE: 9811-008-99
CURRENT APPLICATION NUMBER: US 09/281,717A
CURRENT FILING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: US 60/079,956
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Rushmer, Peter
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR
TITLE OF INVENTION: COACTIVAOR BINDING
TITLE OF INVENTION: WS-999
CURRENT APPLICATION NUMBER: US/09/281,717A
CURRENT APPLICATION NUMBER: US/09/281,717A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

29.5%; Score 31; DB 5; Length 12;
Best Local Similarity 60.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/079,956
PRIOR FILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 35, Application US/09281717A; GENERAL INFORMATION:
APPLICANT: Baxter, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 6, Application US/09830693B
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn version 3.0
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                             (without alignments)
44.130 Million cell updates/sec
                                                                                                                                                                          February 12, 2003, 10:45:03 ; Search time 11 Seconds
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/PUB.WBW_PUB.pep:*

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10: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

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14: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     140259 segs, 25548876 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                             OM protein - protein search, using sw model
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105
1 PPLSQETFSDLWKLLPENG 19
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Gapop 10.0 , Gapext 0.5
                         Copyright
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                                                                                                                                                                                                                                                                                                    Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
                                                                                                                                                                                                                                                                                                                                                                 Sequence:
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		dŧ			SOPPERATES		
Result		Query					
No.	Score	Match	Match Length DB	DB	QI	Description	
н	66	94.3	18	10	US-09-214-371-74	Semience 74 Appl	
7	92	87.6	19	10	US-09-732-384-7		
m	90	85.7	19	12	US-10-155-059-13	Semience 13. April	
4	89	84.8	19	10	US-09-214-371-1		
ហ	77	73.3	15	10	US-09-732-384-6		
9	74	70.5	16	12	US-10-155-059-23	Semience 23. April	
7	99	62.9	12	10	US-09-214-371-17	Semience 17. Appl	
ω	65	61.9	15	10	US-09-950-692-6		
თ	9	57.1	18	12	US-10-155-059-22	Semience 22 Appl	
10	59	56.2	16	10	US-09-214-371-39	Semience 39. April	
11	57	54.3	14	6	US-10-024-123-7	Sequence 7. Appli	
12	55	52.4	12	10	US-09-214-371-7	Sequence 7. Appli	
13	54	51.4	12	10	US-09-214-371-24	Sequence 24. Appl	
14	54	51.4	16	10	US-09-214-371-41	Semience 41 Appl	
15	23	50.5	16	10	US-09-214-371-40	. 4	
16	49	46.7	12	10	US-09-214-371-25	Segmence 25. Appl	
17	49	46.7	12	10	US-09-214-371-26	Sequence 26 Appl	
18	48	45.7	14	10	US-09-214-371-33	33,	
19	48	45.7	14	10	US-09-214-371-34	Sequence 34, Appl	

Sequence 17, Appl Sequence 3, Appli Sequence 21, Appl Sequence 20, Appl	37, 1	35, 36, 2, 7	Sequence 13, Appl Sequence 22, Appl Sequence 46, Appl Sequence 47, Appl Sequence 52, Appl Sequence 52, Appl	Sequence 31, Appl Sequence 33, Appl Sequence 15, Appl Sequence 18, Appl Sequence 5, Appl
12 US-10-155-059-17 10 US-09-019-679-3 11 12 US-10-155-059-21 10 US-09-214-371-20 12 US-10-155-059-24	5 10 US-09-214-371-83 5 10 US-09-732-384-4 5 12 US-10-155-059-1 9 10 US-09-214-371-38	10 US-09-214-371-35 10 US-09-214-371-36 5 12 US-10-155-059-2 9 US-09-840-277-66 7 10 US-09-920-552-110	100 100 100 100	10 US-09-281-71 10 US-09-281-71 10 US-09-281-71 9 US-10-024-123 10 US-09-732-38
444.8 44.8 41.9 11.9 12.04		34.3 30.5 30.5 10 30.5 12 12 12 12 12 12 12 12 12 12 12 12 12	299.55 299.55 299.55 299.55 299.55	29.5 29.5 29.5 12 29.5 14 27.6
20 47 21 44 22 44 23 44 24 42	25 26 27 27 28 36 29 36	30 31 32 32 33 34 32	35 36 37 37 38 39 40	41 31 42 31 44 31 44 55 29

#### ALIGNMENTS

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REBULT.

RESULT.

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REQUENCE 74, Application US/09214371B

Sequence 74, Application US/09214371B

Sequence 74, Application Description

Sequence 7, Application

APPLICANT: Bottger, Volker

APPLICANT: Bottger, Angelica

APPLICANT: Garcia-Encherick

APPLICANT: Garcia-Encherick

PRIOR APPLICANT: Garcia-Encherick

APPLICANT: Garcia-Encherick

CURRENT FLING DATE: 1999-03-26

FILER REFERENCE: 4-199-03-26

FILER REFERENCE: 4-199-03-26

FRICK FLING DATE: 1997-07-04

NUMBER OF SEQ. ID NOS: 83

SOFTWARE: PRIOR APPLICATION NUMBER: PCT/EP97/03549

FRICK FLING DATE: 1997-07-04

NUMBER OF SEQ. ID NOS: 83

SOFTWARE: PRIOR APPLICATION NUMBER: PCT/EP97/03549

FRICK FLING DATE: 1090-03-26

FRICK FLING DATE: 1090-03-26

FRICK FLING DATE: 1090-03-26

SEQ. ID NO 74

FRICK FLING DATE: 1090-03-26

FRICK FLING DATE: 1090-03-26

SEQ. ID NO 74

FRICK FLING DATE: 1090-03-26

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TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                          1 PPLSQETFSDLWKLLPEN 18
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US-10-155-059-13
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US-09-732-384-6
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US-09-214-371-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match 87.6%; Score 92; DB 10; Length 19; Best Local Similarity 100.0%; Pred. No. 6.6e-08; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kaelin, William
APPLICANT: Tost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES
ANTIBODIES
GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR PILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 13, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 617-345-1300
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INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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COMPUTER READABLE FORM:
MEDIUM TYPE: Disket
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COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-155-059-13
                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-732-384-7
                                                                                                                                                                                                                                                             SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Cheme, Patrick
APPLICANT: Cheme, Patrick
APPLICANT: Cheme, Patrick
APPLICANT: Cheme, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20337/AA/PCT;
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR PLING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Description of Artificial Sequence:peptide US-09-214-371-1
                                                                                                        Ouery Match 85.7%; Score 90; DB 12; Length 19; Best Local Similarity 88.9%; Pred. No. 1.3e-07; Matches 16; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match

84.8%; Score 89; DB 10; Length 19;

Best Local Similarity 94.1%; Pred. No. 1.9e-07;

Matches 16; Conservative 1; Mismatches 0; Indels
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APPLICANT: Yuan, Zhi.Min
APPLICANT: Yuan, Zhi.Min
APPLICANT: Gu, JiJie
IITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURENT APPLICATION NUMBER: US/09/732,384
CURENT FILING DATE: 2000-12-07
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 6
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09214371B
Patent No. US20010018511A1
GENERAL INFORMATION:
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Query Match 62.9
Best Local Similarity 100.
Matches 12; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 PPLSQETFSDLWK 13
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US-09-950-692-6
                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                       Gaps
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OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
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70.5%; Score 74; DB 12; Length 16;
Best Local Similarity 81.2%; Pred. No. 2.8e-05;
Matches 13; Conservative 1; Mismatches 2; Indels
                                                             Length 15;
                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: TRM Compatible
COMPUTER: TRM Compatible
CORPATING SYSTEM: Windows
SOFTWARE: FastSED for Windows Version 2.0b
CURRITATION DATA:
PRICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INCRMATION:
                                                         73.3%; Score 77; DB 10; L
100.0%; Pred. No. 9.4e-06;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 23:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Elsenetein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENEL/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 17, Application US/09214371B
                                                                                                                                                                                                                                                                                          Sequence 23, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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                                                                                 Best Local Similarity 100.0
Matches 14; Conservative
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                                                                                                                                                 5 QETFSDLWKLLPEN 18
                                                                                                                                                                                         1 OETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: Boston
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        ; US-09-732-384-6
                                                                    Query Match
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Patent No. US2020106701A1

Patent No. US2020106701A1

APPLICANT: Goueli, Said A

TITLE OF INVENTION: Quantitation of Individual Protein Kinase Activity
TITLE OF INVENTION: Quantitation of Individual Protein Kinase Activity

TITLE OF INVENTION: Quantitation of Individual Protein Kinase Activity

FILE REFERENCE: Kinase

CURRENT FILING DATE: 2010-06-27

PRIOR PELING DATE: 1994-03-04

NUMBER OF SEQ ID NOS: 8

SOFTWARE: PatentIN Ver. 2.0

SEQ ID NO 6

LENGTH: 15
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GENERAL INCOMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION UNMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
CURRENT FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; ; OTHER INFORMATION: polypeptide
US-09-950-692-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61.9%; Score 65; DB 10; Length 15; 84.6%; Pred. No. 0.0006; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62.9%; Score 66; DB 10; Length 12; 100.0%; Pred. No. 0.00033; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 22, Application US/10155059
; Patent No. US20020147173A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 11; Conservative
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Gaps

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FEATURE: OTHER INFORMATION: Description of Artificial Sequence: peptide
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                                                                                                                                                                                                  LOCATION: (1)

OTHER INFORMATION: X = Biotin-Ser
NAME/KEY: VARLANT
LOCATION: (16)

US-09-214-371-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/10024123
Publication No. US20030022263A1
GENERAL INFORMATION:
APPLICANT: Kastan, Michael
APPLICANT: Canman, Christine
                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 11; Conservative
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Best Local Similarity
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                                                                                                                                                                         NAME/KEY: VARIANT
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US-09-214-371-7
                   SEQ ID NO 39
LENGTH: 16
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LENGTH: 14
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                                                             NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE ANTIBODIES
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APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Pickeley, Stephen
APPLICANT: Pickeley, Stephen
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Bree; Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of PS3 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: 05/99/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      57.1%; Score 60; DB 12; Length 18; 76.9%; Pred. No. 0.0041; ative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                           CURRENT INEXELLE
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASESEQ for Windows
SOFTWARE: FASESEQ for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION OF CURROWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/ABORT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOWNUNICATION INFORMATION:
JOST, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 22:
                                                                                                                                                                 ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                          STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 39, Application US/09214371B Patent No. US20010018511A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 18 amino acids
                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 22: SEQUENCE CHARACTERISTICS:
                                                                                                      NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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Best Local Similarity 76.99
Matches 10; Conservative
                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                                                                                                                                   CITY: Boston
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APPLICANT: Canman, Christine
APPLICANT: Kim, Seong-Tae
APPLICANT: Kim, Seong-Tae
APPLICANT: Kim, Seong-Tae
APPLICANT: Lim, Dae-Sik
APPLICANT: St. Jude-Children's Research Hospital
TITLE OF INVENTION: ATM Kinase Modulation for Screening and
TITLE OF INVENTION: Therapies
FILE REFERENCE: 2427/F142
CURRENT APPLICATION NUMBER: US/10/024,123
CURRENT APPLICATION NUMBER: 09/400,653
FRIOR PILING DATE: 1999-09-21
FRIOR PILING DATE: 1999-09-21
FRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 49
SOFTWARE FASISEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   54.3%; Score 57; DB 9; Length 14; 100.0%; Pred. No. 0.0089; Live 0; Mismatches 0; Indels
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Fatent No. US20010018511A1
GENERAL INFORMATION:
APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Gree, Patrick
APPLICANT: Furet, Furet,
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US-09-214-371-40
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APPLICANT:
APPLICANT:
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APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Ghee, Patrick
APPLICANT: Gracia-Echeverria, Carlos
APPLICANT: Huret, Bascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION UNBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence:peptide US-09-214-371-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 51.4%; Score 54; DB 10; Length 12; Local Similarity 100.0%; Pred. No. 0.021; les 10; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                 Query Match
52.4%; Score 55; DB 10; Length 12;
Best Local Similarity 83.3%; Pred. No. 0.015;
Matches 10; Conservative 0; Mismatches 2; Indels
             CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 PRIOR APPLICATION NUMBER: PCT/EP97/03549 PRIOR FILING DATE: 1997-07-04 NUMBER OF SEQ ID NOS: 83 SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 24, Application US/09214371B
; Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (1)
OTHER INFORMATION: X = Ac-Gln
NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
FILE REFERENCE: 4-20937/A/PCT
                                                                                                                                                                                                                                                                                                                                                                                                             5 QETFSDLWKLLP 16
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SEQ ID NO 24
                                                                                                                                                SEQ ID NO 7
LENGTH: 12
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                                                                                                                                                                                     Inhibitors of the Interaction of P53 and MDM2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 51.4%; Score 54; DB 10; Length 16; Best Local Similarity 90.9%; Pred. No. 0.029; Matches 10; Conservative 0; Mismatches 1; Indels
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CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EE97/03549
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SEQ ID NO 40
LENGTH: 16
       APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Ruret, Pascal
TITLE OF INVENTION: Inhibitors of the Interac
FILE REPERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 41
LENGTH: 16
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 40, Application US/09214371B ; Patent No. US20010018511A1
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LOCATION: (1)
OTHER INFORMATION: X = Biotin-Ser
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OTHER INFORMATION: X = Pro-NH2
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COTHER INFORMATION: X = Pro-NH2
US-09-214-371-41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: VARIANT
LOCATION: (1)
OTHER INFORMATION: Biotin-Ser
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APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhik
Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 QETFSDLWKLL 15
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Query Match 50.5%; Score 53; DB 10; Length 16;
Best Local Similarity 90.9%; Pred. No. 0.041;
Matches 10; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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Oy 5 QETFSDLWKLL 15 | | | | | | | | | | Db 5 QPTFSDLWKLL 15 Search completed: February 12, 2003, 10:50:08 Job time : 12 secs

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p53 homologue TIP
p53 N-terminal pep
p53/RB interaction
Human p53 peptide
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Peptide #1. Unide
p53 DR 3a motif bi
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Immunodominant epi
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p53 homologue TIP
                                                                                                                                        February 12, 2003, 10:36:07; Search time 34 Seconds (without alignments)
74.464 Million cell updates/sec
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| SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                         908470 segs, 133250620 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMMARIES
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Maximum Match 100%
Listing first 45 summaries
                                                                                           OM protein - protein search, using sw model
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AAW82319
AAW37228
AAW47074
AAY06310
AAK54909
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AAB29157
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1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                               BLOSUM62
Gapop 10.0 , Gapext 0.5
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Match Length DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             seq length: 0
seq length: 19
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Perfect score:
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Maximum DB
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                                                                                                                                                Run on:
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binding pide 9-25. #3. Unid#7. Unidated pept#11. Unid#11.	#4. #5. is II erac #6.	Ligand #1. Uniden Peptide #2. Unide Human p53 wild-typ Human oncogenic pr Mdm/hdm antagonist Mdm/hdm antagonist	r sast	dig of the contract of the con
AAG897 AAY452 AAB291 AAB291 ABB055 AAB291	AAB2916 AAB2916 AAY9900 AAY5779 AAB2916 AAB2916	AAB2 AAB2 AAW3 AAW3 AAW3 AAB1	ABB7317 ABB7318 AAW3720 AAR8301 AAY9585 ABB0924 AAG8944	AAW3719 AAB1707 AAB1708 ABB7317 ABB7318 AAW3720 AAW3718 AAW3718
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77 7 4 4 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	, , , , , , , ,	66.5 66.5 66 66 66	0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	60 60 60 60 60 60
1122 1132 1155 1155	2222111 22222	2 2 2 2 3 4 3 4 4 3 4 4 4 4 4 4 4 4 4 4	0 0 1 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	6 8 8 8 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4

# ALIGNMENTS

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p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                                New substance with a mdm2 binding domain and coupling partner -useful for stabilising in cells without an efficient mdm2-mediated degradation pathway
       AAW82321 standard; Peptide; 19 AA.
                                                                                                                                                                                                                                                                                        Disclosure; Fig 1; 52pp; English
                                                                                                                                                         98WO-GB01140.
                                                                                                                                                                          97GB-0008089.
                                          (first entry)
                                                            p53 homologue TIP peptide
                                                                                                                                                                                                                               WPI; 1998-609975/51.
                                                                                                                                                                                            (UYDU-) UNIV DUNDEE
                                           22-FEB-1999
                                                                                                                   WO9847919-A1
                                                                                                                                                        20-APR-1998;
                                                                                                                                                                          22-APR-1997;
                                                                                                                                     29-OCT-1998
                                                                                                   Synthetic.
                          AAW82321;
                                                                                                                                                                                                             Lane DP;
AAW8232
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· 0

Pred. No. 5.7e-10;

Best Local Similarity 100.0%;

This sequence is a peptide homologue of a region of p53 which binds to mama. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells on or overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 unction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53. p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament. AAW82319 standard; Peptide; 19 AA. 98WO-GB01144. 97GB-0008092. 1 PPLSQETFSDLWKLLPENG 19 1 PPLSQETFSDLWKLLPENG 19 22-FEB-1999 (first entry) p53 homologue TIP peptide. Best Local Similarity 100.0 Matches 19; Conservative WPI; 1998-609932/51. (UYDU-) UNIV DUNDEE. 19 AA; WO9847525-A1. 20-APR-1998; 22-APR-1997; 29-OCT-1998. Synthetic. Sequence AAW82319; Query Match Lane DP; 셤 ò

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Gaps ö

Length 19; Indels

100.0%; Score 105; DB 19; 100.0%; Pred. No. 5.7e-10; tive 0; Mismatches 0;

This sequence is a peptide homologue of a region of p53 which binds to mand. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells of out overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 unction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53. New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2 Disclosure; Fig 1; 52pp; English.

Sequence 19 AA;

Query Match •

100.0%; Score 105; DB 19; Length 19;

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This represents a p53 N-terminal peptide fragment used in an Blisa TIP assay for analysing the interaction between human oncogenic protein MDM2 and p53. The invention provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated capptosis in tumour cells comprising a wild-type p53 and non-elevated complex and to identify and design inhibitors of MDM2, p53 binding. They can also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia conteractions, especially tumours and viral infections. The peptides can interactions, especially tumours and viral infections. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and caccumulation in normal cells. The peptides which mimic the MDM2 binding content in p53, have a significantly greater blocking activity compared content with wild-type p53.
                                                                                                                                                                                                                                                                                                   MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa TIP assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral
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 Indels
                                                                                                                                                                                                                                                                    p53 N-terminal peptide fragment for Elisa TIP assay.
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100.0%; Pred. No. 4.9e-09;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Furet P,
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Boettger V, Chene P,
Lane D, Picksley S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 34; 45pp; English.
                                                                                                                                                            AAW37228 standard; peptide; 18 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97GB-0007041.
96GB-0014197.
                                   1 PPLSQETFSDLWKLLPENG 19
                                                                    1 PPLSOETFSDLWKLLPENG 19
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                                                                                                                                                                                                                                    20-JUL-1998 (first entry)
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Les 18; Conservative
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-100996/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NOVS ) NOVARTIS AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hochkeppel H,
                                                                                                                                                                                                                                                                                                                                                                                               WO9801467-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-1997;
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JAN-1998.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boettger A,
     19;
                                                                                                                                                                                                AAW37228;
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Synthetic

AAW47074;

AAW47074 RESULT

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wound healing; graft rejection; reperfusion injury;
yrocardial infarction; stroke; traumatic brain injury;
neurodegenerative disease; ischaemia; toxaemia; infection; AIDS;
hepatitis; breast cancer; ovarian cancer; colon cancer; diagnosis;
                                                                                                                        regulatory protein; apoptosis; neoplasia; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This synthetic peptide corresponds to amino acids 13-27 of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New p53 regulatory protein (RB18A) useful as, e.g. sources of and primers to detect the transcription rate and abundance of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR54909 standard; peptide; 15 AA.
                                                                                  Human p53 peptide (aa13-27).
                                                                                                                                                                                                                                                                                                                                                                                                98WO-EP08560.
                                                                                                                                                                                                                                                                                                                                                                                                                                         97EP-0403051.
                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 77.1
Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-395177/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA in lymphocytes
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                                                                                                                                                                                                                                  therapy; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                       15-DEC-1997;
                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                  14-DEC-1998;
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                                         06-SEP-1999
                                                                                                                                                                                                                                                                                                                                                         24-JUN-1999.
                                                                                                                           RB18A; p53
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  AAY06310;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Frade R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a cell enters apoptosis. Apoptosis is prevented if interaction occurs. The interaction is between regions 1-71 or 290-393 of p53 and regions 379-928 of Rb. The invention provides methods to identify compounds able to modulate interaction or binding between p53 and Rb protein. The method and detecting interaction/binding between them. These inhibitory compounds are used in human or veterinary medicine to modulate p53 and rb human or veterinary medicine to modulate p53 activity and processes, specifically for inducing apoptosis in tumour expression of these modulators by gene therapy methods is also contemplated. Other activities that can be affected are transcription repression, G1 arrest, DNA repair, homologous recombination and stabilise p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a p53 protein and a retinoblastoma (RB) protein. This peptide corresponds to residues 11-29 of human p53 (AAW47079). The interaction between p53 and Rb is found to be critical for determining whether or not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This peptide fragment of p53 is an inhibitor of the interaction between
                                                                                                                                                                                                                            p53/RB interaction inhibiting peptide 6 (residues 11-29 of human p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Identifying compounds that modulate interaction of p53 and Rb protein - or those that bind to Rb protein, used to induce apoptosis, specifically for treatment of tumours
                                                                                                                                                                                                                                                                     Retinoblastoma gene; RB; p53 protein; interaction; inhibitor; tumour; apoptosis; modulator; medicine; veterinary; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         92.4%; Score 97; DB 18; Length 19; 94.4%; Pred. No. 1.1e-08; ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CANC-) CANCER RES CAMPAIGN TECHNOLOGY,
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                                                                                                   AAW47074 standard; peptide; 19 AA.
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1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                   Retinoblastoma
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                                                                                                                                                                                   19-MAY-1998
                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
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probes RB18A

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Gaps
p53. It was used to raise anti-peptide p53.1 and anti-RB18A antibodies in rabbit. RB18A (see AAY06309) is a novel p53 regulatory protein of the invention. The antibodies were used in immunoscreenings that led to the isolation of a cDNA clone (see AAX59124) encoding human RB18A. The invention provides methods and compositions for the diagnostic and therapeutic applications of RB18A, in particular for the diagnosis, prevention or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
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                                                                                                                                                                                                                                                              77.1%; Score 81; DB 20; Length 15; 100.0%; Pred. No. 3.2e-06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunodominant epitope from p53 N-terminal.
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Sequence Query Match

Matches

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AAY06310 ID AAY063 XX

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0; Gaps

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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AARS4907-RS4921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              73.3%; Score 77; DB 17; Length 15; 100.0%; Pred. No. 1.4e-05; tive 0; Mismatches 0; Indels
                                                                                                                                       Length 15;
                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                           p53 protein binding motif corresp. to residues 16-30.
                                                                                                                                    73.3%; Score 77; DB 15; I
100.0%; Pred. No. 1.4e-05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                       AAR89914 standard; peptide; 15 AA.
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                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 14; Conservative
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es 14; Conserv
                                                                                                         15 AA;
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20-JUL-1994;
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                                                                          conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                             New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New immuno:dominant epitope(s) of protein p53 - for detecting and monitóring antibodies indicative of cancer and precancerous
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                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Pred. No. --
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                                                                                                                                                                                                                                                       Claim 5; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 42; 62pp; French.
                                                                                                                                    Legros Y, Lubin R, Soussi T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Legros Y, Lubin R, Soussi T;
                                            93WO-FR01082.
                                                                          92FR-0013110.
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                                                                                                       (EURO-) LAB EUROBIO SA.
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nes 14; Conserv
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                                            02-NOV-1993;
                                                                          02-NOV-1992;
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               11-MAY-1994.
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vaccine; epitope; cytostatic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL;
cytotoxic T lymphocyte; T cell; immune response; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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                                                                                                                                                                                       Fork head associated; FHA; domain; transcriptional control;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         73.3%; Score 77; DB 21; L. 100.0%; Pred. No. 1.4e-05; iive 0; Mismatches 0;
                                                                                                                                                                                                       DNA replication; DNA repair; cell cycle control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAG89500 standard; Peptide; 15 AA.
                                                                 AAB29157 standard; Peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Fig 2; 92pp; English.
                                                                                                                                                                                                                                                                                                                              17-MAR-2000; 2000WO-GB01024.
                                                                                                                                                                                                                                                                                                                                                            19-MAR-1999; 99GB-0006432.
28-JUN-1999; 99GB-0015075.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Jackson SP, Durocher D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 PPLSQETFSDLWKL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-664872/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15 AA;
                                                                                                                                                                                                                                                                    WO200057184-A2
                                                                                                                                                                                                                                     Unidentified.
                                                                                                                                                                                                                                                                                                                                                                            28-JUN-1999;
                                                                                                                                                                                                                                                                                                  28-SEP-2000.
                                                                                                                             02-FEB-2001
                                                                                                                                                             Peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAG89500;
                                                                                              AAB29157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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                                      RESULT 9
                                                     AAB29157
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the present inventation describes isolated prepared reas epitopes (1). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (1); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids, (3) a vaccine composition comprising (II), a unit dose of a peptide with less than 50 contiguous amino acids with 100% identity to the native peptide sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II) and (5) an isolated nucleic acid encoding (II). (I) has cytostatic activity and can be used in vaccines. The vaccine composition is useful for treating or preventing cancer. (I) and (II) are useful as diagnostic agents and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine turther, to modify the composition of the epitopes enhances the immunogenicity. The possible pathological side effects caused by intrinsic biological activity, are eliminated. Add89363 to AdG89777 represent annot acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes isolated prepared P53 epitopes (I). Also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL; cytotoxic T lymphocyte; T cell; immune response; tumour; cancer; vaccine; epitope; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Celis E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chesnut R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fikes J, Sette A, Sidney J, Southwood S,
Keogh E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 113; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAG89730 standard; Peptide; 15 AA.
                                                                                                                                                                                                                                           11-DEC-2000; 2000WO-US33629.
                                                                                                                                                                                                                                                                                                                  99US-0458297.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       p53 DR3 binding peptide #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 PPLSOETFSDLWKL 15
                                                                                                                                                                                                                                                                                                                                                                                     (EPIM-) EPIMMUNE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-381493/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 present invention.
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                                                                                                 WO200141788-A1.
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Homo sapiens.
                                                                                                                                                                                                                                                                                                                  10-DEC-1999;
                                                                                                                                                                        14-JUN-2001.
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                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              are useful as diagnostic geners and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine composition of (I). The ability to combine selected epitopes and further, to modify the composition of the epitopes enhances the immunogenicity. The possible pathological side effects caused by infectious agents or whole protein antigens, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Immunisation, immunogenicity enhancer; rabies virus strain ERA; immunòdominant T-helper cell determinant; nucleoprotein; antibody; post-translational modification; p53; cancer; diagnosis.
                                                                                                                                                                                                                 Fikes J, Sette A, Sidney J, Southwood S, Chesnut R, Celis E, Keogh B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                 Example 5; Page 131; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY45227 standard; peptide; 17 AA.
                                                                                                                  11-DEC-2000; 2000WO-US33629.
                                                                                                                                                   99US-0458297.
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                                                                                                                                                                                EPIM-) EPIMMUNE INC.
                                                                                                                                                                                                                                                                    WPI; 2001-381493/40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present invention.
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                                                   WO200141788-A1.
     Homo sapiens.
                                                                                                                                                  10-DEC-1999;
                                                                                  14-JUN-2001.
                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY45227;
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The present invention describes a polypeptide comprising a portion of a by3 comprising a post-translationally modified amino acid residue where the modification is a phosphorylation, glycosylation or prenylation, and an immunogenicity enhancer. The polypeptide is used to detect anti-p53 autoantibodies in a human patient. The antibody can be used to determine the post-translational modification state of a p53 protein. These are used especially in the diagnosis of cancer. The present sequence represents a p53 peptide used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                             Methods of detecting post-translational modification level of p53, useful for diagnosis of cancer \, -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fork head associated, FHA, domain, transcriptional control, DNA replication; DNA repair, cell cycle control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   73.3%; Score 77; DB 20; Length 17; 100.0%; Pred. No. 1.6e-05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                       Thurin M, Hoffman R;
                                                                                                                                                                                                                                                                                                                    Disclosure; Page 45; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB29159 standard; Peptide; 15 AA.
                                                                                                           98US-0077512
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99GB-0006432.
99GB-0015075.
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es 14; Conservative
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                                                                                                                                                 (WIST-) WISTAR INST.
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                                                                                                                                                                                   Ertl H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 AA;
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                                                                       11-MAR-1999;
W09946574-A2
                                                                                                             11-MAR-1998;
                                   16-SEP-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
                                                                                                                                                                                     Otvos L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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15 AA;

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                               The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assays and screening methods based on direct interaction between FHA
                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                               70.5%; Score 74; DB 21; Length 15; 92.9%; Pred. No. 4.2e-05; ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB29163 standard; Peptide; 15 AA.
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Disclosure, Fig 2; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAR-2000; 2000WO-GB01024.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Durocher D;
                                                                                                                                                                                                                                                                                                                                                              13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                   2 PPLAQETFSDLWKL 15
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                                                                                                                                                                                                                                                                                                                                          Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide #7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   binding -
                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                   Ubiquitin dependent proteolysis modulation; cdc4 phospho design motif; CDP motif; cytostatic; nootropic; antiproliferative; cell proliferation; growth; differentiation; cancer; neurodegenerative disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             that targets molecules for ubsquirin dependent processives. (C) have cytostatic, nootropic and antiproliferative activity. Also described is a method for the treatment of a disease or condition where affected cells have a defective protein, comprising administering (C) to promote degradation of the target protein in cells by ubiquitin dependent proteolysis. (C) can also be used for modulating the proliferation, growth and/or differentiation of cells. (C) can be used to modulating the proliferation, differentiation of cells. (C) can be used to modulate ubiquitin dependent proteolysis or cell proliferation, growth and or differentiation of cells. (C) is useful in the treatment of cancers and neurodegenerative disorders as well as spinal degeneration. The present sequence represents a biotinylated peptide which is used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a cdc4 phospho design (CPD) motif, (C),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New Cdc4 Phospho Design motif that targets molecules for ubiquitin dependent proteolysis, is useful for the modulation of cell
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Score 74; DB 21; Length 15;
Pred. No. 4.2e-05;
1; Mismatches 0; Indels
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100.0%; Pred. No. 5.6e-05;
tive 0; Mismatches 0;
                                   1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proliferation i.e. cancer treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 2; Page 47; 83pp; English.
                                                                                                                                                                                      ABB05528 standard; Peptide; 14 AA.
                                                                                                                                                                                                                                                                                     Biotinylated peptide SEQ ID NO:13.
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   70.5%;
92.9%;
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24-JAN-2001; 2001US-263774P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAY-2001; 2001WO-CA00632.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        from the present invention.
                                                                                                                                                                                                                                                         (first entry)
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Matches 13; Conservative
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                                                                    1 PPLSQETFSDLWKL 14
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 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Modified-site
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                                      Matches
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Search completed: February 12, 2003, 10:43:48 Job time : 35 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:42:32; Search time 28 Seconds (without alignments) 139.818 Million cell updates/sec Run on:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Title:

Perfect score:

Sequence:

Gapop 10.0 , Gapext 0.5 **BLOSUM62** Scoring table:

671580 seqs, 206047115 residues Searched:

Total number of hits satisfying chosen parameters:

5270

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SPTREMBL 21:\* Database :

sp archea:\*
sp\_bacteria:\*
sp\_fungi:\*
sp\_human:\*
sp\_invertebrate:\*
sp\_mammal:\*
sp\_phage:\*
sp\_phage:\*
sp\_plant:\*
sp\_plant:\*
sp\_rodent:\*
sp\_vortebrate:\*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

### Q9bgg8 sorex arane 077538 bos taurus 07538 bos taurus 020401 bomo sapien 82326 pisum sativ Q9qvs7 mus sp. lac Q9qvs7 mus sp. lac Q91513 homo sapien Q81960 influenzavi Q81093 paracoccus P82674 bos taurus Q84129 influenzavi Q22064 dioscorea t Q81965 influenzavi Q81955 influenzavi Q81955 mus musculu P83003 entamoeba h Description P82326 Q9QVS7 Q9UJZ3 Q83960 Q9F0P3 Q83965 Q9MJQ5 Q9ESU5 P82674 Q84129 P83003 Q9BGG8 O77538 Q9XNQ1 Q9UEL0 110 110 110 110 110 110 Query Match Length DB Score Result

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7 8 8 8 7 7 7 7 8 8 8 8 7 7 7 7 8 8 8 8	17 15 072017 18 13 013167 19 6 019107 7 8 095945 9 4 0943X3 13 11 091XP1
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# ALIGNMENTS

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SEQUENCE FROM N.A.
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SEQUENCE
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Jiang H., Lucy M.C.;
"Identification of 5' untranslated variants of bovine growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jiang H., Lucy M.C.; "Identification of 5' untranslated variants of bovine growth hormone
                                                                                                                                                                         Larkin D., Serov O., Zhdanova N.;
"Mapping of five genes from human chromosome 17 to chromosome hn of
the common shrew (Sorex araneus).";
Acta Theriol. (Warsz) 85:143-146 (2000).
EMBL; AF314827; AAK13419.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=BRBED HOLSTEIN;
BRDLINE=99175163; PubMed=10075683;
Jiang H., Okamura C.S., Lucy M.C.;
"Isolation and characterization of a novel promoter for the bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
MCBI_TaxID=9913;
                                                                                                       Sorex araneus (Eurasian common shrew) (European shrew).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Insectivora, Soricidae, Soricinae, Sorex.
                                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                                                                                                    . Match 29.5%; Score 31; DB 6; Length 16; Local Similarity 60.0%; Pred. No. 3.4e+02; les 6; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jiang H., Lucy M.C.; "Bobine GH receptor 5' UTR variants."; Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases. EMBL, AF036297; AAC33315.2; "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (DEC-1997) to the BMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                               SEQUENCE 16 AA; 1951 MW; 775186E3FE5F52E2 CRC64;
                                            01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
Thyroid hormone receptor alpha (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1998 (TrEMBLrel. 08, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth hormone receptor gene.";
J. Biol. Chem. 274:7893-7900(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Growth hormone receptor (Fragment).
                        PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                         PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bos taurus (Bovine).
                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSD 10
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                                                                                                                                               NCBI_TaxID=42254;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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Parasitiformes; Ixodida; Ixodidae; Boophilus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27.6%; Score 29; DB 8; Length 17; 75.0%; Pred. No. 7.6e+02; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                Length 14;
                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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                                                                                                                                                                                                                                                                                   14 14 14
14 AA; 1544 MW; 5A45D458FF3A9E27 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UUN-2001 (TrEMBLrel. 17, Last annotation update)
FAS antigen (CDS antigen) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
Cytochrome oxidase subunit 2 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                Score 30; DB 6; I
Pred. No. 4.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                      1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boophilus microplus (Cattle tick).
EMBL, AF085281, AAD22521.1; --
EMBL, AF036290, AAC33308.1; --
EMBL, AF036292, AAC33310.1; --
EMBL, AF036293, AAC33310.1; --
EMBL, AF036293, AAC33311.1; --
EMBL, AF036295, AAC33312.1; --
EMBL, AF036295, AAC33312.1; --
EMBL, AF036295, AAC33313.1; --
EMBL, AF036296, AAC33314.1; --
EMBL, AF326349; AAC33314.1; --
                                                                                                                                                                                                                                                                                                                                                                Similarity 83.3%; 5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FAS antigen (CD95 antigen)
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les 6; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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Q9UJZ3;
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                                                                                            "Transcription stimulation of the Fas-encoding gene by nuclear factor for interleukin-6 expression upon influenza virus infection."; J. Biol. Chem. 270:18007-18012(1995).
                                                                                                                                                                                                                                                                                                                                                                                     Muschen M., Re D., Jungnickel B., Diehl V., Rajewsky K., Kuppers R.; "Somatic mutations of the CD95 gene in human B cells as a side-effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUN-2000 (TrEMBLrel. 14, Created)
01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
01-JUN-2000 (TrEMBLrel. 15, Last annotation update)
Unknown protein from 2D-gage of thylakoid lumen (SpOTIO7) (Fragment).
Pisum sativum (Garden pea).
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
eurosids 1; Fabales; Fabaceae; Papilionoideae; Vicieae; Pisum.
NCBL TaxID=3888;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                      SEQUENCE FROM N.A.
Muschen M., Re D., Brauninger A., Wolf J., Hansmann M.L., Diehl V.,
Kuppers R., Rajewsky K.;
"Somatic mutations of the CD95 gene in Hodgkin- and Reed-Sternberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pelifier J. B., Friso G., Kalume D.E., Roepstorff P., Nilsson F., Adamska I., van Wijk K.J.;

"Proteomics of the chloroplast: systematic identification and targeting analysis of lumenal and peripheral thylakoid proteins.";

Plant Cell 12:319-341(2000).

-I. SUBCELLULAR LOCATION: CHLOROPLAST THYLAKOID MEMBRANE LUMEN.

-I. DEVELOPMENTAL STRAE: UNFOLDED AND FULLY DEVELOPED LEAVES.

-I. MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN CHOOLOGIST IS, 60, ITS MW IS: 18.3 KDA.

CHOROPLAST IT, 14, 14, 14, 14, 14, 14
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0
                    MEDLINE-95355401; PubMed-7543095;
Wada N., Matsumura M., Ohba Y., Kobayashi N., Takizawa T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24.8%; Score 26; DB 10; Length 14; 83.3%; Pred. No. 1.9e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25.7%; Score 27; DB 4; Length 11; 66.7%; Pred. No. 1e+03; 1; Indels ive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             of the germinal center reaction.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; D31968; BAZ08501;
EMBL; AJ279012; CAC35540.1;
EMBL; AJ279013; CAC35540.1;
AJ279013; CAC35540.1;
AJ279013; CAC35540.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE, SUBCELLULAR LOCATION, AND DEVELOPMENTAL STAGE.
STRAIN-CV. DE GRACE; TISSUE-LEAF;
MEDLINE-20181728; PubMed-10715320;
                                                                                                                                                                                                                                                                                                               Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE 11 AA; 1256 MW; 0802145E2731B370 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                               "Molecular analysis of four lactate dehydrogenase-A mutants in the
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NAR-2002 (TrEMBLrel. 20, Last annotation update)
11-NAR-2002 (TrEMBLrel. 20, Last annotation update)
11-NAR-2004 (H212), non-structural protein (Seg 8), cooh terminus of nsl (Fragment).
                                                                                                                                                                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Sandulache R., Pretsch W., Chatterjee B., Gimbel W., Graw J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
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"Human oviduct glycoprotein promoter sequence.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF189710; AAF01065.1; -.
NON TER
SEQÜENCE 18 AA; 2201 MW; 5E8FD91EA210E516 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                            mouse.";
Mamm. Genome 5:777-780(1994).
SEQUENCE 17 AA; 1982 MW; E941E0A3F2477D45 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                             Last sequence update)
Last annotation update)
17 AA.
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                                                01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last seq
01-DBC-2001 (TrEMBLrel. 19, Last ann
Lactate dehydrogenase-A (Fragment).
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Best Local Similarity 30.8
Matches 4; Conservative
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Matches 4; Conservative
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PRELIMINARY;
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Bos taurus (Bovine).
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                                                                       NCBI_TaxID=9913;
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Bedzyk L.A., Escudero K.W., Gill R.E., Griffin K.J., Frerman F.E.;
"Cloning, sequencing, and expression of the genes encoding subunits of
Paracoccus denitrificans electron transfer flavoprotein.";
J. Biol. Chem. 268:20211-20217(1993).
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MEDLINE=21101849; PubMed=11160087;
Fales L., Kryszak L., Zeiletra-Ryalls J.;
"Control of hemA Expression in Rhodobacter sphaeroides 2.4.1: Effect of a Transposon Insertion in the hbdA Gene.";
J. Bacteriol. 183:1568-1576(2001).
                                                                                                                                                                                                         Gaps
                                                                        Parvin J.D., Young J.F., Palese P., "nonsense mutations affecting the lengths of the ns1 nonstructural proteins of influenza a virus isolates."; Virology 128:512-517(1983).
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Bacteria; Proteobacteria; alpha subdivision; Rhodobacter group;
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Influenzavirus A.
Viruses; ssRNA negative-strand viruses; Orthomyxoviridae;
Influenza A viruses.
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                                                                                                                                                              SEQUENCE 16 AA; 1811 MW; 23F434B5AEEFAD93 CRC64;
                                                                                                                                                                                                                                                                                                                                     01-MAR-2001 (TrEMBLrel. 16, Last sequence update) 01-DEC-2001 (TrEMBLrel. 19, Last annotation update) Electron transfer flavoprotein subunit A (Fragment)
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01-OCT-2000 (TrEMBLrel. 15, Created)
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                                                     SEQUENCE FROM N.A.
MEDLINE=83303830; PubMed=6612993;
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InterPro; IPR000256; Flu NS1.
Pfam; PF00600; Flu NS1; 1.
NON TER
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Best Local Similarity 50.۰۰
ابر A; Conservative
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nes 4; Conservative
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                                 NCBI_TaxID=11320;
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01-MAR-2001
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 20, Last annotation update)
Influenza ASwine/Wisconsin/OM/2 (HINI), non-structural protein (Seg B), COOH terminus of NSI (Fragment).
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Innesses mutations affecting the lengths of the ns1 nonstructural
profesins of influenza a virus isolates.";
Virology 128:512-517(1983).
Wirology 128:512-517(1983).
InterPro; IPR000256; Flu NS1.
Pfam; PF00600; Flu NS1; I.
                                                                                                        Bukaryota, Merazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Cetartiodactyla, Ruminantia, Pecora, Bovoidea,
Bovidae, Bovinae, Bos.
                                                                                                                                                                                                                                                                                                                                                      "Identification of four proteins from the small subunit of the mammalian mitochondrial ribosome using a proteomics approach."; Submitted (UUL-2000) to the SWISS-PROT data bank.
-i- SUBCELLULAR LOCATION: MITOCHONDRIAL.
-i- SIMILAR LOCATION: MW=974.58; METHOD=ELECTROSPRAY; RANGE=1-8.
-i- SIMILARITY: BELONGS TO THE SSP FAMILY OF RIBOSOMAL PROTEINS.
InterPro; IPRO10851; Ribosomal SSP FAMILY.
PROSITE; PROSESS; RIBOSOMAL SS; PARTIAL.
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
Mitochondrial 28S ribosomal protein S5 (MRP-S5) (Fragments).
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18 18
18 AA, 2127 MW; 199BC913E7E25FAF CRC64;
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                                                                                                                                                                                                                                                         SEQUENCE, AND MASS SPECTROMETRY.
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Local Similarity 57.1%;
les 4; Conservative 5
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NCBI_TaxID=11320;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Influenza A.berkeley/1/68 (H2n2), non-structural protein (Seg 8), cooh terminus of nsl (Fragment).
                                                                                                        Dioscorea tenuipes.
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Dioscoreales, Dioscoreaceae,
                                                                                                                                                                                                                                Terauchi R., Terachi T., Miyashita N.;
"DNA polymorphism aut the Pgi locus of a wild yam, Dioscorea tokoro.";
Genetics 147:1899-1914 (1997).
EMBL; AB006004; BAA23204.1; -.
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Parvin J.D., Young J.F., Palese P.;

Parvin J.D., Young J.F., Palese P.;

Pronsense mutations affecting the lengths of the nsl nonstructural proteins of influenza a virus isolates.";

Virology 128:512-517(1983).

EMBL; K00957; AAA43518.1; -.

InterPro; IPR000256; Flu NSl.

Pfam; PF00600; Flu_NSl; Ī.

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Pred. No. 3.8e+03;
2; Mismatches 7; Indels
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Viruses; seRNA negative-strand viruses; Orthomyxoviridae;
Influenza A viruses.
VCBI_TaxID=11320;
                                                                                                                                                                                                                                                                                                    SEQUENCE 19 AA; 2235 MW; E10F1743F4263EAE CRC64;
                                     01-JAN'1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Phosphoglucose isomerase (EC 5.3.1.9) (Fragment).
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MEDLINE=98072503; PubMed=9409845;
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30.8%;
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NCBI_TaxID=64476;
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Saguez C., Lecellier G., Koll F.;
Saguez C., Lecellier G., Koll F.;
Intronic GIY-YIG endonuclease gene in the mitochondrial genome of Padospora curvicolla: evidence for mobility.";
Nucleic Acids Res. 28:1299-1306 (2000).
EMBL; Ad49984; CAB72447.1;
InterPro: IPRO00179; Cyt. b. b6.
PROSITE; PS00133; CYTOCHROWE_B_Q0; UNKNOWN_1.
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                                                                                                           Eukaryota; Pungi; Ascomycota; Pezizomycotina; Sordariomycetes; Sordariales; Lasiosphaeriaceae; Podospora.
                                                                                                                                                                                                                                                                                                                                                                                                    22.9%; Score 24; DB 8; Length 10; 80.0%; Pred. No. 2.8e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 10 AA; 1200 MW; C31A223B437B0772 CRC64;
            01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UN-2001 (TrEMBLrel. 17, Last annotation update)
Cytochrome b (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Matches 4; Conservative
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                                                                                     Podospora curvicolla.
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:36:28; Search time 29 Seconds (without alignments) 27.174 Million cell updates/sec Run on:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 segs, 41476328 residues Searched:

1015 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt\_40:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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SUMMARIES		ΩĪ	HTF HELZE	AKHX LOCMI	NP1 LYMST	NP2_LYMST	NP4_LYMST	NP5_LYMST	IBP4 PIG	UR2_SCYCA	HTF2 PERAM	HTF2_CARMO	URE1 MORMO	ACT6_SOYBN	CPAX_BOVIN	AKH_TABAT	HTF1_PERAM	HTF_TABAT	NP3_LYMST	PH4_PERAM	PCG6_PACGO	AL22_HORSE	IRBP_CAVPO	PCG7_PACGO	AKHG_GRYBI	CLP_THICU	HTF_TENMO	HTF_NAUCI	URA7_HUMAN	CHI1_PEA	LEC1 PSOSC	PC20_BRANA	ALL1_CALVO	FIBA ANTAM	HI70_RAT
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FKB7_PINPS HS11_PINPS	RBS PHYPA AKH LIBAU	RPCH_PANBO	SLAP_BACTG	POOC PSEFL	Q2OA COMTE	NSK2 SARBU	UC34 MAIZE	ARCA_STRP5	LEC3_PSOSC
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# ALIGNMENTS

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Pancrustacea; Lepidoptera; Helicoverpa	T. lc lc (T (TII	CARBOXYLIC ACID B9D1 CRC64; B1; Length 10, 3e+02; s 3; Indels	10 AA.  pdate) lupdate)  .update, bulata; Pancrustacea; bolata; Orthoptera; Oedipodinae; Locusta.
) Pancr Lepid ; Heli	., Bird eptide } pokinet: 988). NEUROPI MOLYMPH OF INSEK	BOXYLIC 1 CRC64 ; Leng 02; 3; I	
10 AA. date) update ). ulata; ygota;	tr B.A., neuropepl d adipol 350 (198 KS ARE NUTH HEMO)	B A5.	10 AA. odate) update) ulata; eroidea;
PRT; 1 ) quence upd quetion u e2-HRIn a; Mandibu Endoptery	Fraser; a net of a net and si344-355344-3E IN THE HEMOLYN	PYRROLIDONE CARBC AMIDATION: 8E70367865A5B9D1 Score 24; DB 1; Pred: No. 5.3e+02; ; Mismatches	
PRT; 10 AA. ed) sequence update) annotation update) (HeZ-HRTH). (HoZ-HRTH). oda; Mandibulata; a; Endopterygota; idae; Heliothinae;	15690; C.T., es D.K cture halose un. 15 AEMIC EHALOS IN THE		PRT; 10 AA sed) sed) sequence update) annotation updat ry locust). ry locust). ry locust). oda; Mandibulata a; Orthopteroide crididae; Oedipo
	TISSUE=Corpora cardiaca; TISSUE=Corpora cardiaca; MEDINE=88326324; PubMed=3415690; MEDINE=88326324; PubMed=3415690; Taffe H., Raina A.K., Railey C.T., Fraser B., Tseng C.M., Zhang Y.S., Hayes D.K.; "Isolation and primary structure of a neuroi Hellothis zea with hypertrehalosemic of an encorpic and primary structure of a neuroi Biochem. Biophys. Res. Commun. 155:344:350(-1-FUNCTION: HYDERTREHALOSSEMIC FACTORS ARI ELEVATE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH -1-SIMILARITY: BELONGS TO THE AKH / HRTH / InterPro; IPRO02047; AKH.	KH; 1. tion. 10 1096 MW; 22.9%; 62.5%; ative	eat st atc rop rer
STANDARD; 1. 15, Cr 1. 28, La 11. 28, La mic hormo Corn earw Zoa; Arth octa; Neop	ardiac ; Pubh A.K., g Y.S. rimary th hyg th hyg PERTRI LEVEL LEVEL LEVEL BELON 571.	0256; AKH; 1.  Amidation.  10  0 AA; 1096 M  larity 62.5  Conservative  W 12  W 8	STANDARD; 1. 39, Cr 1. 39, La 1. 41, La 1. 41, La cria (Migr cria (Migr cria (Migr cria (Migr cridoidea
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ELZE FIGATE STANDARD; FIG53; 01-AUG-1990 (Rel. 15, Creat 01-FEB-1994 (Rel. 28, Last Hypertrehalosaemic hormone Hypertrehalosaemic hormone Haliothis zea (Corn earworn Fisheryota; Metazoa; Arthrop Insecta; Perrygota; Noctuoidea; Noctuoi	SEQUENCE. TISSUE=Corpora cardiaca; MEDLINE=88326324; Pubmed Jaffe H., Raina A.K., Ri Tseng C.M., Zhang Y.S., i "Isolation and primary s Holothis zea with hyper. Blochem. Blophys. Res. C! FUNCTION: HYPERTREHA! FUNCTION: THE LEVEL OF THE MAJOR CARBOHYDRA -! SIMILARITY: BELONGS' -! SIMILARITY: BELONGS' PTR; A31571; A31571; A31571.	PROSITE; PS00256; AKH; 1 Neuropeptide; Amidation. MOD_RES 10 10 SEQUENCE 10 AA; 1096 ry Match 22. t Local Similarity 62. ches 5; Conservative 5 QETFSDLW 12	T 2 LOCMI STANIANA PB1626, 30-MAY-2000 (Rel. 3; 15-JUN-2002 (Rel. 4) Peptide hormone. Locusta migratoria Eukaryota, Metazoa; Insecta; Pterygota; Acridomorpha; Acrid
HTF HELZE ID HTF HELZE AC P16353; DT 01-RTB- DT 01-RTB- DT 01-RTB- DT 01-RTB- DT HTP- DT 01-RTB- DT	11] SUUSUCE TISSUE=C MEDILNE= MEDILNE= TISSUE C.: TISSUE C.: HIJOTH HIJOTH BEOCHEM. -! FUNC -! FUNC -! FUNC -! SIMI PIR, SIMI	PROSITE; Neuropeping MOD_RES MOD_RES SEQUENCE Query Match Matches SAME SECTION	RESULT 2  AKHX LOCMI  D AKHX LOCMI  AC P81626,  DT 30-MAY-2006  DT 15-JUN-2005  DE Peptide hor  SC Eukaryota;  CC Insecta; Processor Acridomorph  OC Acridomorph
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora; Lymnaeidae; Lymnaea.
NCBL TaxID=6523;
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Bukaryota, Metazoa, Mollusca, Gastropoda, Pulmonata, Basommatophora,
                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=93238777; PubMed=8477756;
Johnsen A.H., Rehfeld J.F.;
JimmanFamides, a new family of neuropeptides from the pond snail,
Lymnae stagnalis. Clue to cholecystokinin immunoreactivity in
invertebrates?";
                 Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; 832471; S32471.
Neuropeptide; Amidation.
MOD_RES.
13 13 13
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4; Conservative 1; Mismatches 3; Indels
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                                                                                                          PYRROLIDONE CARBOXYLIC ACID.

        MOD RES
        13
        AMIDATION.

        SEQÜENCE
        13 AA;
        1519 MW;
        9CA07BA3F5D5B455 CRC64;

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MOD_RES 10 10 AMIDATION.
SEQUENCE 10 AA; 1222 MW; 81BFF67AB415B9D1 CRC64;
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01-JUL-1993 (Rel. 26, Last annotation update)
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NCBI_TaxID=6523;
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Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
Lymnaeidae; Lymnaea.
NCBI_TaxID=6523;
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Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
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TISSUB-Ganglion;
MEDLINE-93238777; PubMed-8477756;
Anohneen A.H., Rehfeld J.F.;
"LymnaDFanides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
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Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.

NOUR PEGE 13 AMIDATION.
                                                                                                                                                                                                                                                                                  invertebrates?";
Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARINY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; 532472; 532472.
Nouropeptide; Amidation.
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SEQÜENCE 13 AA; 1503 MW; 9CA07BBB56D5B455 CRC64;
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01-UUL-1993 (Rel. 26, Last annotation update)
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MEDLINE=93238777; PubMed=8477756;
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8 FSD-LWK 13
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HTF2_PERAM
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                         Johnsen A.H., Rehfeld J.F.;
"LymnaDFamides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92109718; PubMed=1722398;
Coleman M.B., Pan Y.-C.E., Etherton T.D.;
Identification and MI2-terminal amino acid sequence of three insulin-like growth factor-binding proteins in porcine serum.";
Biochem. Biophys. Res. Commun. 181:1131-1136 (1991).
-! FUNCTION: IGFS-BINDING PROTEINS PROLONG THE HALF-LIFE OF THE IGNOMENTAND HAVE BEEN SHOWN TO EITHER INHIBIT OR STIMULATE THE GROWTH PROMOTING EFFECTS OF THE IGFS ON CELL CULTURE. THEY ALTER THE INTERFACTION OF IGFS WITH THEIR CELL SURFACE RECEPTORS.
-! SUBCELLULAR LOCATION: Secreted.
-! SIMILARITY: CONTAINNS 1 THYROGLOBULIN TYPE-I DOMAIN.
-! SIMILARITY: EBLONGS TO THE INSULIN-LIKE GROWTH FACTOR BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                            Sus scrofa (Pig).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Cetartiodactyla; Suina; Suidae; Sus.
NCBI_TaxID=9823;
                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                  01-MAR-1992 (Rel. 21, Created)
01-MAR-1992 (Rel. 21, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Insulin-like growth factor binding protein 4 (IGFBP-4) (IBP-4)
(IGF-binding protein 4) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .
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                                                                         Eur. J. Biochem. 213:875-879(1993).
-1- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; S32475; S32475.
Neuropeptide, Amidation.
MOD RES
SEQUENCE 13 A3, 1476 MW; 9CA07BBBS6DSASS CRC64:
                                                                                                                                                                           21.0%; Score 22; DB 1; Length 13; 50.0%; Pred. No. 1.40+03; vative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 22; DB 1; Length 16;
Pred. No. 1.8e+03;
1; Mismatches 1; Indels
                                                                                                                                                 9CA07BBB56D5A5A5 CRC64;
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InterPro; IPR0000667; Insl_gro_fac_pr.
InterPro; IPR000016; Thyroglobulin_1.

PROSITE; PS00222; IGF BINDING; PARTIAL.

PROSITE; PS00484; THYROGLOBULIN_1; PARTIAL.

Growth factor_binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Mismatches
              MEDLINE=93238777; PubMed=8477756;
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13 AA; 1476 MW;
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1es 4; Conservative
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                                                                  invertebrates?";
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7 PPPSEE 12
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ID IBP4 PIG
AC P24854;
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UR2_SCYCA
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12 AA.

STANDARD;

UR2\_SCYCA P35490;

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MEDLINE-84298179; PubMed-6591205;
Scarborough R.M., Jamieson G.C., Kalish F., Kramer S.J., McEnroe G.A.,
Miller C.A., Schooley D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Blatta orientalis (Oriental cockroach).

Bukaryota, Metazoa, Arthropoda, Mandibulata; Pancrustacea, Hexapoda;
Insecta; Persygota; Neoptera; Orthopteroidea, Dictyoptera; Blattaria,
Blattoidea, Blattidee, Periplaneta.

NCBI_TaxID=6978, 7539, 6976;
                                                                                                                                                                                                                                                                                                 MEDLINE=92319231; PubMed=1620290;

Conlon J.M., O'Harte F., Smith D.D., Balment R.J., Hazon N.;

"Purification and characterization of urotensin II and parvalbumin from an elasmobranch fish, Scyliorhinus canicula (common dogfish).";

Neuroendocrinology 55:230-235(1992).

-! FUNCTION: HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A CORTICOTROPIN-RELEASING FACTOR. PROBABLY INVOLVED IN SMOOTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Hypertrehalosaemic factor II (Neuropeptide M-II) (Periplanetin CC-2)
Periplaneta americana (American cockroach),
Leptinotarsa decemineata (Colorado potato beetle), and
01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-201 (Rel. 40, Last annotation update)
Urotensin II (U-II) (UII)
Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
NCBI_TaxID=7830;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Structures of two cockroach neuropeptides assigned by fast atom bombardment mass spectrometry."; Biochem. Biophys. Res. Commun. 124:350-358(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPECIES=P.amerioana;
MEDLINE=85046530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M., Cook J.C., Hemling M.E.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "Isolation and primary structure of two peptides with cardioacceleratory and hyperglycemic activity from the corpora cardiaca of Periplaneta americana mericana proc. Natl. Acad. Sci. U.S.A. 81:5575-5579 (1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 20.5%; Score 21.5; DB 1; Length 12; Similarity 71.4%; Pred. No. 1.6e+03; 5; Conservative 0; Mismatches 1; Indels
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12 Aa; 1526 MW; 804729F9D579CEBA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MUSCLE STIMULATION.
-!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.
InterPro; IPR001483; Urotensin II.
PRO2083; Urotensin II; II.
PROSITE; PS00984; UROTENSIN_II; I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
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                                                                                                                                                                                                                                         MEDLINE=90253659; PubMed=2340112;
Gaede G., Rinehart K.L. Jr.;
"Primary structures of hypertrehalosaemic neuropeptides isolated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Extatosoma tiaratum (Stick insect).

Eukaryota; Metazoa; Arthopda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Pterygota; Neoptera; Orthopteroidea; Phasmatodea;
Heteronemiidae; Carausius.
                                                                                                                                                                                                                                                                                                      the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insec Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry.";

Biol. Chem. Hoppe-Seyler 371:345-354(1990).

-!- FUNCTION: HYPERTREHALOSAEMIC FACTORS ARE NEUROPEPTIDES THAT ELEVALE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH (TREHALOSE IS THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).

-!- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
            SPECIES=1.decemlineata; TISSUE=Corpora cardiaca;
MEDLINE=90160053; PubMed=2576128;
Gaede G., Kellner R.;
"The metabolic neuropeptides of the corpus cardiacum from the potato
beetle and the American cockroach are identical.";
Peptides 10:1287-1289(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUL-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
30-WAY-2000 (Rel. 39, Last annotation update)
Hypertrehalosaemic factor II (HTF-II) (HRTH-II) (Hypertrehalosaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gade G., Rinchart K.L. Jr.; Gade G., Rinchart K.L. Jr.; Primary structure of the hypertrehalosaemic factor II from the corpus cardiacum of the Indian stick insect, Carausius morosus, determined by fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
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                                                                                                                                                                                                                   SPECIES=B.orientalis; TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuropeptide II).
Carausius morosus (Indian stick insect), and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES=C.morosus; TISSUB=Corpora cardiaca;
MEDLINE=87157103; PubMed=3828078;
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Interpro; IPR002047; AKH.
PROSITE; PS00256; AKH: 1.
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S08996; S08996.
B44960; B44960.
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CARBOHYDRATE-LINKAGE SITE.

CARBOHYDRATE-LINKAGE SITE.

KEDINE-93129188; PubMed=1482345;

KEDLINE-93129188; PubMed=1482345;

KEDLINE-93129188; PubMed=1482345;

Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;

Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;

Ta stick finsect corpus cardiacum.";

Biochem. Biophys. Res. Commun. 189:1303-1309 (1992).

LIBIOCHEM. BIOPHYS. Res. Commun. 189:1303-1309 (1992).

CIPETATE THE LEVEL OF TREALORS ARE NEWOPEPTIDES THAT ELVERTORY THE HEMOLYMPH (TREHALOSE IS THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).

CIPETANALOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).

CIPETANALOR CARBOHYDRATE IN THE MEMOLYMPH OF INSECTS.
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01.AUG-1990 (Rel. 15, Last sequence update)
15.Unv.2002 (Rel. 41, Last annotation update)
Urease alpha subunit (EC 3.5.1.5) (Urea amidohydrolase) (Urease 63 kDa subunit) (Fragment).
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Gromphadorhina portentosa, Blattella germanica and Blatta orientalis
                           and of the stick insect Exxatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 371:345-354 (1990).
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Hu L.-T., Nicholson E.B., Jones B.D., Lynch M.J., Mobley H.L.T.;
Hu L.-T., Nicholson E.B., Jones B.D., Lynch M.J., Mobley H.L.T.;
"Morganella morganii urease: purification, characterization, and
isolation of gene sequences.";
J. Bacteriol. 172:3073-3080(1990).
-!- CATALYTIC ACTIVITY: Urea + H(2)0 = CO(2) + 2 NH(3).
-!- COPACTOR: Binds 2 nickel ions per subunit (Potential).
-!- SUBUNIT: (ALPHA, BETA, GAMMA) (3) (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE UREASE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morganella morganii (Proteus morganii).
Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 21; DB 1; Length 10;
Pred. No. 1.5e+03;
1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C-LINKED (MAN) (PROBABLE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15 AA; 1650 MW; 09E27AA54241687B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9B9036745771A9D1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20.0%; Score 21; DB 1; L 25.0%; Pred. No. 2.4e+03; tive 5; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro, IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation; Glycoprotein.
MOD_RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00145; UREASE 2; PARTIAL. PROSITE; PS01120; UREASE_1; PARTIAL. Hydrolase; Metal-binding; Nickel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; A35389; A35389.
InterPro; IPR001924; UreaseA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20.0%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 AA; 1164 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
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Matches 3; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QLTFTPNW 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Morganella.
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SEQUENCE
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VARIANT
NON TER
SEQUENCE
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AKH_TABAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation use by mon-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            o;
                                                                                                                                                                                Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae; Glycine.
NCBI_TaxID=3847;
                                                                                                                                                                                                                                                                               Pearson L., Meagher R.B.;
"Diverse soybean actin transcripts contain a large intron in the 5'
untranslated leader: structural similarity to vertebrate muscle actin
                                                                                                                                                                                                                                                                                                                                                                 IN ALL EUKARYOTIC CELLS.
FUNCTION: BSSENTIAL COMPONENT OF CELL CYTOSKELETON; PLAYS AN IMPORTANT ROLE IN CYTOPLASMIC STREAMING, CELL SHAPE DETERMINATION, CELL DIVISION, ORGANELLE MOVEMENT AND EXTENSION GROWTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                               Flant Mol. Biol. 14:513-526(1990).
-!- FUNCTION: ACTINS ARE HIGHLY CONSERVED PROTEINS THAT ARE INVOLVED IN VARIOUS TYPES OF CELL MOTILITY AND ARE UBIQUITOUSLY EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SUBCELLULAR LOCATION: CYCOPLAGMIC.
-!- MISCELLANEOUS: THERE ARE AT LEAST 16 ACTIN GENES IN SOYBEAN.
-!- SIMILARITY: BELONGS TO THE ACTIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 AA; 1749 MW; 38F4970D4BEFF3C3 CRC64;
                                                                                                         01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
                                                                                      17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00022; actin; 1. PRTIAL. PROSITE; PS00406; ACTINS 1; PARTIAL. PROSITE; PS00432; ACTINS 2; PARTIAL. PROSITE; PS01132; ACTINS ACT LIKE; PARTIAL. Structural protein; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CPAX BOVIN STANDARD; PRT; 18 AA. P22779; 01-AUG-1991 (Rel. 19, Created) 01-AUG-1991 (Rel. 19, Last sequence update)
                                                                                                                                                                                                                                                        STRAIN=cv. Wayne;
MEDLINE=91346640; PubMed=2102831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL, X17119; CAA34979.1; -. PIR; S15754; S15754. InterPro; IPR004001; Actin. InterPro; IPR004000; Actin_like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4, Conservative
                                                                                    STANDARD;
                                                                                                                                                                       Glycine max (Soybean)
1 PPLSQETFSDLW 12
                         1 POISROEYGGLF 12
                                                                                                                                               Actin 6 (Fragment).
                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
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                                                                                    SOYBN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQÜENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                   ACT6 SOY
P15986;
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                                                          RESULT 12
ACT6_SOYBN
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TISSUB=Corpora cardiaca;

XX MEDLINE=80046/78; Pubmed=2813385;
XX MEDLINE=80046/78; Pubmed=2813385;
Adfe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Vogel V.W., Zhang Y.-S., Hayes D.K.;
Vogel V.W., Zhang Y.-S., Hayes D.K.;
The Vogel V.W., Stang Y.-S., Research of hormones with adipokinetic and hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of horse of hypotrehalosemic activity isolated from the corpora cardiaca of horse of horse
                                                                                                                                                                                                                                                                                                                                                                                                                   15-DEC-1998 (Rel. 37, Last annotation update)
Cytochrome P450 2A (OLF2) (Olfactive) (P52) (EC 1.14.14.1) (Fragment).
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Insecta, Pterygota, Neoptera, Endopterygota, Diptera, Brachycera,
Tabanomorpha, Tabanidae, Tabanus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PIR; A35704; A35704.

PIRs; A35704; A35704.

PROSITE; PS00086; CYTOCHROME_P450.

PROSIGNE; PROBLES; CYTOCHROME_P450; PARTIAL.

Oxidoreductase; Monocoxygenase; Blectron transport; Membrane; Heme; Microsome; Endoplasmic reticulum; Olfaction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- CATALYTIC ACTIVITY: RH + reduced flavoprotein + O(2) = ROH + oxidized flavoprotein + H(2)O.
-!- SUBCELLULAR LOCATION: Membrane-bound. Endoplasmic reticulum.
-!- SIMILARITY: BELONGS TO THE CYTOCHROME P450 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Adipokinetic hormone (AKH) (Dipteran corpora cardiaca factor I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20.0%; Score 21; DB 1; Length 18; 36.4%; Pred. No. 2.9e+03; ive 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 AA; 2058 MW; F80746F76CCD77FF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACIDS, AND XENOBIOTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 36.41,
A; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
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18
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us-09-403-440a-2.closed.rsp

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SPECIES=P.americana;
MEDLINE=84298179; PubMed=6591205;
Scarborough R.M., Jamieson G.C., Kalish F., Kramer S.J., McEnroe G.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Primary structures of the cockroaches Leucophaea maderae, the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry.", Biol. Chem. Hoppe-Seyler 371:345-354 (1990).

-: FUNCTION: HYBERTEHALOSAEMIC FACTORS ARE NEUROPEPTIDES THAT ELEVART THE LEVEL OF TREHALOSE IN THE HEMOLYWPH (TREHALOSE IS 'THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Hypertrehalosaemic factor I (Neuropeptide M.)
(PeA-CAH-I) (LeD-CC-I) (Hypertrehalosaemic neuropeptide I).
Periplaneta americana (American cockroach),
Leptinotaras decemilineata (Colorado potato beetle), and
Blatta orientalis (Oriental cockroach).
Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Perrygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria;
Blattoidea; Blattidae; Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPECIES=L.decemlineata; TISSUE=Corpora cardiaca;
MEDLINE=90160053; PubMed=2576128;
Gaede G., Kellner R.;
"The metabolic neuropeptides of the corpus cardiacum from the potato
beelle and the American cockroach are identical.";
Peptides 10:1287-1289(1989).
                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Structures of two cockroach neuropeptides assigned by fast atom bombardment mass spectrometry."; Biochem. Biophys. Res. Commun. 124:350-358(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPECIES=P.americana;
MEDIJINE=85946530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M., Cook J.C., Hemling M.E.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Miller C.A., Schooley D.A., "Isolation and primary structure of two peptides with cardioacceleratory and hyperglycemic activity from the corpora
                                          Neuropeptide; Amidation; Flight.

MOD RES 1 1 1 PYRROLIDONE CARBOXYLIC ACID.

MOD_RES 8 AMIDATION
                                                                                                                                                                                                                                                Query Match
19.0%; Score 20; DB 1; Length 8;
Best Local Similarity 50.0%; Pred. No. 1.1e+05;
Matches 4; Conservative 1; Mismatches 3; Indels

        MOD_RES
        8
        AMIDATION.

        SEQUENCE
        8 AA; 949 MW; 86786771A9D1A736 CRC64;

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Proc. Natl. Acad. Sci. U.S.A. 81:5575-5579(1984).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
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InterPro; IPR002047; AKH
                               PS00256; AKH; 1
                                                                                                                                                                                                                                                                                                                                                                                                       5 QETFSDLW 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 QLTFTPGW 8
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                                  PROSITE;
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Gaps
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                                                                                                                                                                                                                      4; Indels
                                                                                                                                                                                         Length 8;
-i- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
                                                                                                                                                         SEQUENCE 8 AA; 991 MW; 86745775B9C452D6 CRC64;
                                                                                                                                                                                       Query Match
19.0%; Score 20; DB 1; I
Best Local Similarity 50.0%; Pred. No. 1.1e+05;
Matches 4; Conservative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                    Search completed: February 12, 2003, 10:44:24
                                                                                                                                            AMIDATION
             PIR, A05169, A05169.
PIR, S08995, S08995.
PIR, A49823, A49823.
PIR, A44960, A44960.
InterPro, IPR002047, AKH.
PROSITE, PS00256, AKH, 1.
                                                                                                             Neuropeptide; Amidation.
MOD RES 1 1
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:42:53; Search time 46 Seconds Run on:

(without alignments) 39.708 Million cell updates/sec

US-09-403-440A-2 Perfect score:

1 PPLSQETFSDLWKLLPENG 19 Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283224 seqs, 96134422 residues Searched:

3433 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

PIR\_73:\* Database :

pirl:\* pir2:\* pir3:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		de			SUMMARIES	
Result		Ouerv				
No.	Score	Match	Length	DB	ID	Description
-	34	32.4	19	2	B61409	
7	26	24.8	18	7	A59396	
m	25	23.8	14	~	A35105	him p I - Inaumeto
4	25	23.8	15	~	PA0110	intportierical proce
Ŋ	24	22.9		'n	A31571	translation elonga
9	24	22.9	-	η (	S41747	nypertrehalosemic/
7	23	21.9		~	S11545	chaperonin 10 homo
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57	21		10	N	JC1416	9 .
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27	21	20.0	10	0	\$53789	III d
28	21	20.0	11	~	3300	peptide Pec
29	21	0	1.	۰,	200	214K excantigen (v
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30 21 20.0 13 2 B35245 31 21 20.0 13 2 B56845 33 21 20.0 13 2 B56864 34 21 20.0 13 2 B56864 35 21 20.0 15 2 A45189 36 21 20.0 16 2 B58503 37 21 20.0 17 2 B15754 38 20.0 17 2 B15754 39 20.5 19.5 13 2 P41627 41 20.5 19.5 18 2 I52623 42 20 19.0 8 2 A49823 44 20 19.0 8 2 A449823 45 20 19.0 8 2 A449823 46 20.5 19.0 8 2 A449823 47 20 19.0 8 2 A449823 48 20 19.0 8 2 A449823	histone H1.c - mouhistone H1a - mous dipeptidyl-peptida urease (EC 3.5.1.5 7 alpha-hydroxy-4-superoxide dismuta actin 6 - soybean cytochrome p450 ol TCR delta chain VCSteine proteinasigh chain V-D-D-Trhypothetical proteinasighypothetical proteinalosemical proteinasighypothetical proteinasighypothetic hormonal proteinasighypothetic proteinasighypothetic hormonal proteinasighypothetic proteinasighypothetical
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	0 H W W W W W W W W W W W W W W W W W W

## ALIGNMENTS

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Genome polyprotein (clone L3/S2) - Skalica virus (fragment)
C;Species: Skalica virus
C;Species: 19-Mar-1997 #sequence_revision 19-Dec-1997 #text_change 17-Nov-2000
C;Accession: B61409
R;Guirakhoo, F.; Heinz, F.X.; Mandl, C.W.; Holzmann, H.; Kunz, C.; Gresikova, M.
J. Gen. Virol. 72, 333-338, 1991
A;Title: The relationship between the flaviviruses Skalica and Langat as revealed by month?Reference number. A61409; MUID:91132129; PMID:1847173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                 A/Status: preliminary; not compared with conceptual translation A/Molecule type: genomic RNA A/Residues: 1-19 <GUI> C/Superfamily: yellow fever virus genome polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

32.4%; Score 34; DB 2; Length 19;
Best Local Similarity 41.7%; Pred. No. 81;
Matches 5; Conservative 3; Mismatches 4; Indels
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B61409
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2 PPVTYRTGTDCW 13 q

RESULT 2 A59396

That is a remainded possible procession of the procession of the procession of the procession and characterization of a major allergen of the procession has been considered to the Protein Sequence Database, September 2001 A; Reference number: A59396

A;Status: preliminary A;Molecule type: protein A;Residues: 1-18 <MON>

A; Experimental source: L5

A; Note: IgE-binding protein, major allergen

0; Gaps 24.8%; Score 26; DB 2; Length 18; 38.5%; Pred. No. 1.28+03; tive 3; Mismatches 5; Indels 5; Conservative Best Local Similarity Matches 5; Conserv Query Match

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6 ETFSDLWKLLPEN 18 ò

2 ETYSDKYDTIDVN 14 a

RESULT 3

beta-D-galactosida

Genetics:

Matches

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C; Accession: A24244
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Holman, G.M.; Wagner, R.M.; Ridgwa Biochem. Biophys. Res. Commun. 135, 622-628, 1986
A;Title: Isolation and primary structure of a peptide from the corpora cardiaca of Heli A;Reference number: A24244; MUID:86186794; PMID:3964263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Accession: A24244
A;Molecule type: protein
A;Residues: 1-9 c.ARF?
A;Superfamily: adippkinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone, neuropeptide; Fyroglutami
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R;Gaede, G.; Wilps, H.; Kellner, R.
Biochem, J. 269, 309-313, 1990
A;Title: Isolation and structure of a novel charged member of the red-pigment-concentral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adipokinetic hormone - bollworm
N;Alternate names: Hez-AKH
C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Species: 31-Mar-1988 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                adipokinetic hormone - nestling-sucking blowfly
C;Species: Protophormia terraenovae (nestling-sucking blowfly)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 31-Oct-1997
                                                                                                                                                                                                                                                                                        C;Species: Solanum tuberosum (potato)
C;Date: 19-Mar-1997 #Sequence_revision 29-Aug-1997 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                         R;Burt, W.J.E.; Leaver, C.J.
FEBS Lett. 339, 139-141, 1994
A;Title: Identification of a chaperonin-10 homologue in plant mitochondria.
A;Reference number: S41747; MUID:94148071; PMID:7906228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21.9%; Score 23; DB 2; Length 8; 62.5%; Pred. No. 2.8e+05; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 22.9%; Score 24; DB 2; Length 11; Local Similarity 83.3%; Pred. No. 1.5e+03; es 5; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A, Experimental source: mitochondrion C, Keywords: mitochondrion; molecular chaperone
                                                                                                                                                                                                                                                             chaperonin 10 homolog - potato (fragment)
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A;Residues: 1-8 <GAE>
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Best Local Similarity
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A, Residues: 1-11 < BUR>
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| ||| |
1 QLTFSSGW 8
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C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Date: 30-Unn-1989 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
C;Accession: A31571
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Bird, T.G.; Tseng, C.M.; Zhang, Y.S
B;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Bird, T.G.; Tseng, C.M.; Zhang, Y.S
B;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Bird, T.G.; Tseng, C.M.; Zhang, Y.S
A;Title: Isolation and primary structure of a neuropeptide hormone from Heliothis zea wi
A;Reference number: A31571; MUID:88326324; PMID:3415690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Superfeamily: adipokinetic hormone C;Keywords: neuropeptide; pyroglutamid C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid F;I/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;10/Modified site: amidated carboxyl end (Asn) #status experimental
                                              hypothetical protein - Neurospora crassa mitochondrion (fragment)
hypothetical protein - Neurospora crassa
Cipate: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 07-Dec-1999
C;Accession: A35105
C;Accession: A35105
Cell 61, 685-696, 1990
A;Title: A site-specific self-cleavage reaction performed by a novel RNA in neurospora 1A;Reference number: A35105; MUID:90263093; PMID:2160856
A;Accession: A35105
A;Accession: A35
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22.9%; Score 24; DB 2; Length 10;
Best Local Similarity 62.5%; Pred. No. 1.3e+03;
Matches 5; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23.8%; Score 25; DB 2; Length 14; 54.5%; Pred. No. 1.3e+03; tive 0; Mismatches 5; Indels
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R; Kamo, M.; Kawakami, T.; Tsugita, A. submitted to JIPID, March 1995
A; Reference number: PA0109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Keywords: mitochondrion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Residues: 1-14 <SAV>
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hypothetical protein 2 - Arabidopsis thaliana (Species: Arabidopsis thaliana (Mouse-ear cress)
C;Species: Arabidopsis thaliana (Mouse-ear cress)
C;Date: 21-Nov-1993 #sequence_revision 12-May-1995 #text_change 21-Jul-2000
C;Accession: S16324
R;Ruberti, I.; Sessa, G.; Lucchetti, S.; Morelli, G.
R;Ruberti, I.; Sessa, G.; Lucchetti, S.; Morelli, G.
A;Title: A novel class of plant proteins containing a homeodomain with a closely linked A;Reference number: S16323; MUID:91266907; PMID:1675603
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Eur. J. Biochem. 213, 875-879, 1993
A.Title: LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaea stagm
A.Reference number: S32471; MUID:93238777; PMID:8477756
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C.Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
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C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
C;Accession: S32472
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A,Residues: 1-13 «JOH»
A,Cross-references: PIDN:AAB26362.1; PID:g299829
A,Experimental source: ganglia
C;Keywords: amidated carboxyl end, neuropeptide
F;13/Modified site: amidated carboxyl end (Phe) #status predicted
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A/Residues: 1-13 <JOH>
A/Residues: 1-13 <JOH>
A/Residues: 103 <JOH>
A/Cross-references: PIDN:AAB26363.1; PID:g299830
A/Experimental source: ganglia
C/Keywords: amidated carboxyl end; neuropeptide
F/13/Modified site: amidated carboxyl end (Phe) #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21.0%; Score 22; DB 2; Length 8; 80.0%; Pred. No. 2.8e+05; tive 1; Mismatches 0; Indels
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Matches 4; Conservative
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Matches 4; Conserva
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A; Residues: 1-8 < RUB>
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3 YKLLP 7
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PEBS Lett. 298, 33-35, 1992
A;Title: Phosphorylation of cytochrome b6 by the LHC II kinase associated with the cytochrome number: S20410; MUID:92183823; PMID:1544419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Description: is responsible for the regulation of energy distribution between photosys A; Note: does not exhibit redox-controlled activation C; Keywords: chloroplast; light-harvesting complex; membrane-associated complex; phosphoc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Drotein kinase (BC 2.7.1.37) - spinach chloroplast (fragment)
N;Alternate names: LHCII protein kinase
C;Species: chloroplast Spinacia oleracea (spinach)
C;Date: 19-Mar-1997 #sequence_revision 11-Jun-1999 #text_change 11-Jun-1999
C;Accession: S20410
                                                                                                                                                                                                          Gaps
F:1/Modifised site: pyrrolidone carboxylic acid (Gln) #status experimental F:9/Modifised site: amidated carboxyl end (Gly) #status experimental
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                                                                                                                  21.9%; Score 23; DB 2; Length 9; 50.0%; Pred. No. 2.8e+05; Live 1; Mismatches 3; Indels
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                                                                                                                                                    Local Similarity 50.0
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Matches 4; Conserv
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C; Function:
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Cispecies: Lymnaea stagnalis (great pond snail)
Cjacte: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
CjAccession: 832475
RjOhnsen, A.H.; Rehfeld, J.F.
By Dhiochem. 213, 875-879, 1993
A;Title: LymnaPanides, a new family of neuropeptides from the pond snail, Lymnaea stagn A;Reference number: 832471; MUD:93238777; PMID:8477756
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Cypecies: lymnaea stagnalis (great pond snail)
Cybec: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
C;Accession: 832474
C;Accession: 832474
C;Accession: 213, 875-879, 1993
A;Itile: LymnaParanides, a new family of neuropeptides from the pond snail, Lymnaea stagn. A;Reference number: 832471; MUID: 93238777; PMID: 8477756
A;Accession: 832474
A;Accession: 832
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A;Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
P;13/Modified site: amidated carboxyl end (Phe) #status predicted
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21.0%; Score 22; DB 2; Length 13;
Best Local Similarity 50.0%; Pred. No. 3.5e+03;
Matches 4; Conservative 1; Mismatches 3; Indels
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                                                                                                                   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
             GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-245-500A-1

US-08-390-546-1

US-08-390-546-1

US-08-390-5167-1

US-08-390-515A-1

US-08-697-221-19

US-08-697-221-20

US-08-697-221-22

US-08-697-221-22

US-08-697-221-23

US-08-697-221-23
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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seq length: 200000000
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Maximum DB :
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APPLICANT: Kaelin, William
APPLICANT: Abor, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF 7
TITLE OF INVENTION: ANTIBODIES
TITLE OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
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9, Appli
2, Appli
6, Appli
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32, Appli
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Sequence 11,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION DATE:
APPLICATION NUMBER: US/09/091,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Elsensein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCY/DOCKET NUMBER: 30,628
REFERENCY/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEDSOMMUNICATION INFORMATION:
TELEDSOMMUNICATION INFORMATION:
                         US-08-1390-516C-7

US-08-1390-516C-8

US-08-130-516C-9

US-08-130-516C-9

US-08-130-515A-6

US-08-130-515A-7

US-08-130-515A-7

US-08-130-515A-9

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Patent No. 6451979
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101 Federal Street
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LENGTH: 20 amino acid
TYPE: amino acid
STRANDEDNESS: single
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US-09-081-975-8
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usa
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US-09-081-975-16
                                                                   COUNTRY:
STREET:
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GENERAL INFORMATION:

APPLICANT: Kaelin, William

APPLICANT: Jost, Christine

TITLE, OF INVENTION: METHODS OF TREATMENT USING

TITLE, OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE

TITLE OF INVENTION: ANTIBODIES
                                                                                                                                                                                                                                  APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MATIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IEM Compatible
COMPUTER: IEM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastERG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTOMNEY/AGENT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-081-975-11
; Sequence 11, Application US/09081975
; Patent No. 6451979
                                                                                                                                                                 Sequence 10, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 10:
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       1 PPLSOETFSDLWKLLPEN 18
                                            2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSE: Nixon Pea
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US-09-081-975-10
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Sequence 16, Application US/09081975

Patent No. 6451979

GENERAL INFORMATION:
APPLICANT: Realin, William
APPLICANT: Realin, William
TITLE OF INVENTION: METHODIES AND PROTEINS THERETO, AND USES OF THE TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS: 27

ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                            OPERATING SYSTEM: Windows SOFTWARE: FASTEM: Windows SOFTWARE: FASTESQ for Windows Version 2.0b CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/ABENT INFORMATION:
NAME: PICLATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READBLE FORM:
MBDIUM TYPE: Diskette
COMPUTER I IBM Compatible
COMPUTER: IEM COMPATIBLE
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRNT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 474
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 101 Federal Street
CITY: Boston
101 Federal Street
                                                                                                                                                                  COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 617-345-605
TELEFAX: 617-345-1300
                                                                                                                                                Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-081-975-11
                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
                         Boston
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ZIP: 02110
                                                                                               02110
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                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: HOLLE, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 5
CORRESPONDENCES: 5
CORRESPONDENCES: 5
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 G STREET; N.W.
                                                                        Query Match 94.3%; Score 99; DB 1; Length 64; Best Local Similarity 100.0%; Pred. No. 5.9e-09; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94.3%; Score 99; DB 1; Length 64; 100.0%; Pred. No. 5.9e-09; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READBLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,546
FILING DATE: 07-APR 1993
CLASSIFICATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELEPHONE: 202-508-9100
TELEPHONE: 202-508-929
TELENT: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                   RESULT 6
US-08-390-546-1
; Sequence 1, Application US/08390546
; Patent No. 5606044
                                                                                                                                                                              1 PPLSQETFSDLWKLLPEN 18
       CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                            12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-terminal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: WASHILL STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FRAGMENT TYPE: NORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
            ;
US-08-245-500A-1
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APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID B.
APPLICANT: KINZLER, KENDETH W.
APPLICANT: KINZLER, KENDETH W.
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN TITLE OF SEQUENCES: 5
CORRESPONDENCES: 5
CORRESPONDENCES: 5
ADDRESSE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 GETREET, N.W.
                                                                                                                                                                                                                                                                                                                                                                    Query Match 94.3%; Score 99; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 1.6e-09; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/245,500A
FLING DATE: 07-APR-1993
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08245500A
Patent No. 5550023
                                                                                                                                                           INFORMATION FOR SEQ ID NO: 16:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

US-09-081-975-16
               REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47,
TELECOMMUNICATION INPORMATION:
TELEPHONE: 617-345-6054
TELEPAX: 617-345-1300
Eisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 64 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 5
US-08-245-500A-1
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ZIP: 20001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CORRESPONDENCE ADDRESS: ADDRESSE: BANNER, B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
STREET: 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-557-393-1
                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: V
STATE:
                                                                                      STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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94.3%; Score 99; DB 1; Length 64;
Best Local Similarity 100.0%; Pred. No. 5.9e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08557393
Patent NO. 5702903
GENERAL INFORMATION:
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN ITILE OF INVENTION: AMPLIFICATION OF HUMAN TUMORS
NUMBER OF 'SEQUENCES: 5
                                                                                                      APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: KINELER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
WIMBER OF INVENTION: HUMAN TUMORS
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/390,479A
FILING DATE: 02-FEB-1995
CLASSIFICATION: 530
ATYONEN'AGENT INPORMATION:

NAME: REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 32,141
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION S00-929
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                   ADDRESSEE: BANNER & WITCOFF, LTD. STREET: 1001 G STREET, N.W. CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
WIBLIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                               Sequence 1, Application US/08390479A Patent No. 5618921 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 PPLSQETFSDLWKLLPEN 29
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FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 64 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                   STATE: D.C. COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                              20001
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                             US-08-390-479A-1
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US-08-557-393-1
         RESULT 7
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Sequence 1, Application US/08390516C
Patent No. 5708136
GENERAL INFORMATION:
APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: VOGELSTEIN, BERT
ITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATE: US/08/557,393
FILLING DATE: 13-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%; Score 99; DB 1; Le
100.0%; Pred. No. 5.9e-09;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SSEE: BANNER, BIRCH, MCKIE AND BECKETT: 1001 G STREET, N.W. WASHINGTON
E: BANNER, BIRCH, MCKIE AND BECKETT 1001 G STREET, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PPLICATION
PILING DATE: 13-NOV-1995
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/245,500
FILING DATE: 18-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
TELEBHONE: 202-508-9100
TELEBHOME: 202-208-9100
                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                            ZIP: 2001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 18; Conservative
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                                                                                                         WASHINGTON
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APPLICANT: BURELL, MARILEE
APPLICANT: BURELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 G STREET, N.W.
CITY: WASHINGTON
STATE: D.C.
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                   94.3%; Score 99; DB 1; Length 64; 100.0%; Pred. No. 5.9e-09; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 1, Application US/08390515A; Patent No. 5756455
       TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                    N-terminal
                                                                                                 LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 100.8
Matches 18; Conservative
                         TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D.C.
USA
                                                                                                                                                                                                                                  FRAGMENT TYPE: NORIGINAL SOURCE:
                                                                                                                                                                                                                    ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                               US-08-390-517A-1
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US-08-390-515A-1
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Sequence 1, Application US/08390517A;
Fatent No. 57338
GENERAL INFORMATION:
APPLICANT: BURRELL, MARILEE
APPLICANT: KINZETH W.
APPLICANT: KINZETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: ADSTREET: N.W.
CITY: WASHINGTON
STATE: D.C.
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
94.3%; Score 99; DB 1; Length 64;
Best Local Similarity 100.0%; Pred. No. 5.9e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                             CURRENT APPLICATION DATA:

SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/390,516C
FILING DATE: 07-APR.1993
CLASSIFICATION: 530
ATTONEY/AGENT INPORMATION:
NAME: REGISTRATION NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
REFERENCE/POCKET NUMBER: 01107.42798
TELECOMMULCATION INFORMATION:
TELEPHONE: 202-508-920
TELEFAX: 202-508-929
TELEX: 197430 BBMB UT
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 2001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: DAtentil Release #1.0, Version #1.25
SOFTWARE: Patentil Release #1.0, Version #1.25
GURREWT APPLICATION DATE: 07-APR-1993
FLING DATE: 07-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
REFERENCE/POCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-terminal
                                                                                                                                                                                                                                                                                                                         LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTI-SENSE: NO
FRAGMENT TYPE: N
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-390-516C-1
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0; Gaps

0

Gaps 0;

0; Indels

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0; Mismatches
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SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                                                                                                                         HILL, DAVID E.
KINZLER, KENNETH W.
VOGELSTEIN, BERT
                                                                                                                                                                                     Sequence 1, Application US/09170159A; Patent No. 6399755; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 202-508-9100
TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                    APPLICANT: BURRELL, MARILEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT TYPE: N-terminal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12 PPLSQETFSDLWKLLPEN 29
                                      1 PPLSQETFSDLWKLLPEN 18
                                                                        12 PPLSQETFSDLWKLLPEN 29
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: D.C.
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 18; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 20001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-170-159A-1
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US-09-170-159A-1
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                                                                                                                                              Query Match

94.3%; Score 99; DB 1; Length 64;
Best Local Similarity 100.0%; Pred. No. 5.9e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: El POPPY disk
COMPUTER: DEP C. COMPATIBLE
CORRANG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/801,718
FILING DATE: 14-FEB-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/390,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ...UKESSEE: BANNER, BIRCH, MCKIE AND BECKETT STREET: 1001 G STREET, N.W. CITY: WASHINGTON STATE: D.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/390,515
FILING DATE: 07-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEFAX: 202-508-929
INFORMATION FOR SEG ID NO: 1: SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/08801718
Patent No. 5858976
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.3%; S
100.0%;
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CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-terminal
                                                                                                                                                                                                                                       1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                    12 PPLSQETFSDLWKLLPEN 29
                                           ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
      N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: si
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Best Local Similarity
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D.C.
USA
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    FRAGMENT TYPE: DORIGINAL SOURCE:
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                                                                                                            US-08-390-515A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-801-718-1
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0; Gaps
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN HUMAN TUMORS
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: IBM FC compatible
COMPUTER: DEADBLE FOLDOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATE: 13-00c-1998
TILING DATE: 13-0ct-1998
CLASSIFICATION: Cubricoma-
ATTORNEY AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.3%; Score 99; DB 4; Length 64; ilarity 100.0%; Pred. No. 5.9e-09; Conservative 0; Mismatches 0; Indels
                                                                                     CORRESPONDENČE ADDRESS:
ADDRESSES: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 G STREET, N.W.
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 3, Application US/09414436
; Patent No. 6294384
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APPLICANT: Dell'Acqua, Giorgio
APPLICANT: Dell'Acqua, Giorgio
APPLICANT: Datu, Wichael J.
APPLICANT: Mann, Michael J.
APPLICANT: Datu, Victor J.
TITLE DESU, Victor J.
FILE REFERENCE: D53f
CURRENT APPLICATION NUMBER: US/09/414,436
CURRENT APPLICATION NUMBER: US/09/414,436
CURRENT PILING DATE: 1999-10-07
EARLIER APPLICATION NUMBER: 60/103,849
EARLIER FILING DATE: 1999-10-09
SOFTWARE: PALENTH Ver: 2.0
SEQ ID NO 3
LENGTH: 261
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 94.3%; Score 99; DB 4; Length 261; Best Local Similarity 100.0%; Pred. No. 2.8e-08; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 17, Application US/08697221
; Sequence 17, Application US/08697221
; Patent No. 584708:
; Patent No. 584708:
; APPLICANT: Halazonetis, Thanos D.
; TITLE OF INVENTION: Modified p53 Constructs and Uses
; TITLE OF INVENTION: Therefor
; NUMBER OF SEQUENCES: 33
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr., PO Box 457
CITY: Spring House
STATE: Pennsylvania
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 19477
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: ER PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/697,221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/004,802
FILING DATE: 22-SEP-1995
ATTORNEY AGENT INFORMATION:
NAME: KOGROFF, CALHY A.
REGISTRATION NUMBER: 33,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WST64AUSA
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-540-9206
TELEFAX: 215-540-5818
INPORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 363 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 PPLSQETFSDLWKLLPEN 18
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MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-697-221-17
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        Matches
        18;
        Conservative
        0;
        Mismatches
        0;
        Indels
        0;
        Gaps
        0;

        QY
        1
        PPLSQETFSDLWKLLPEN
        18
        Db
        12
        PPLSQETFSDLWKLLPEN
        29

        Search
        completed:
        February
        12, 2003, 10:36:21
        Job time:
        15 secs
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1 PPLSQETFSDLWKLLPEN 18
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Sequence 220, App
Sequence 898, App
Sequence 25, Appl
Sequence 27, Appl
Sequence 27, Appl
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Sequence 21, App
Sequence 27, Appl
Sequence 27, Appl
Sequence 286, App
Sequence 250, App
Sequence 5, App
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Sequence 3, Appli
Sequence 9, Appli
Sequence 4, Appli
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3137, Ap
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Sequence 56463, A
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Sequence 118, App
Sequence 9, Appli
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                                                                       February 12, 2003, 10:35:07; Search time 19 Seconds (without alignments) 83.787 Million cell updates/sec
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Sequence 3
Sequence 3
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1. /ggn2 6/ptodata/1/paa/USO6_NEW COMB.pep:*

2. /ggn2 6/ptodata/1/paa/USO6_NEW COMB.pep:*

3. /ggn2 6/ptodata/1/paa/USO8_NEW COMB.pep:*
        GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-724-676-56463

US-09-724-676-56463

US-09-99-602-24

PCT-USO2-21461-3

US-09-860-24

US-10-274-814-4

US-60-423-552-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-10-218-282-898

US-10-23-128-898

US-10-233-553-28

US-10-233-553-28
                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                         425801 segs, 83787487 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUMMARIES
                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                    OM protein - protein search, using sw model
                                                                                                                                    105
1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                   Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                         US-09-403-440A-2
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Match Length DB
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                                                                                                                                                                       Scoring table:
                                                                                                                                   Perfect score:
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51.5
50
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3470, Ap
6, Appli
77164, A
                                                                                                                                                          17, Appl
1032, Ap
29, Appl
156, App
2160, Ap
3137, Ap
4695, Ap
7, Appli
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77164, A
77166, A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SEQ TYRARE: PatentIn version 3.2
SEQ ID NO 55463
LENGTH: 351
                    PCT-USO2-32851-7
US-10-072-012-607
US-10-072-012-608
US-10-072-012-608
US-10-031-822-49
PCT-USO2-3277-3470
US-10-057-498-3470
US-10-057-498-3470
US-09-724-676-77164
US-09-724-676-77164
US-09-724-676-77166
US-09-724-676-77166
US-10-163-587A-17
US-10-276-781-103
PCT-USO2-32851-29
PCT-USO2-32851-29
PCT-USO2-32851-29
PCT-USO2-32851-29
US-10-264-237-2160
US-10-264-237-2160
US-10-264-237-2160
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100.0%; Pred. No. v.-
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US-09-124-676A-56463
IS-GQUENCE 56463, Application US/09724676A
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                ; Sequence 56463, Application US/09724676; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12 PPLSQETFSDLWKLLPEN 29
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Best Local Similarity 100.0
Matches 18, Conservative
397
434
837
2828
2828
2828
70
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1481
1481
1481
1724
1724
612
617
271
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ORGANISM: Homo sapiens
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0.11400
0.114044
0.005
0.005
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Wills, Ken N.
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                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
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US-10-274-874-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%; Score 99; DB 5; Length 353; 100.0%; Pred. No. 6.3e-08; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: BIXIL PHARMACCULICALS, INC. et al.
TITLE OF INVENTION: SIR2 ACTIVITY
FILE REPERBENCE: 13407-015W01
CURRENT APPLICATION NUMBER: PCT/US02/21461
CURRENT PILING DATE: 2002-07-08
FRICR APPLICATION NUMBER: US 10/190,159
FRICR APPLICATION NUMBER: US 60/303,370
FRICR APPLICATION NUMBER: US 60/303,456
FRICR APPLICATION NUMBER: US 60/303,456
FRICR FILING DATE: 2001-07-06
FRICR FILING DATE: 2001-07-06
NUMBER OF SEQ ID NOS: 24
SOFTWARE: FASSEQ for Windows Version 4.0
                                                                                                                                                                           APPLICANT: Old, Lloyd J.
APPLICANT: Stockert, Elisabeth
APPLICANT: Check, Yao-Tseng
TITLE OF INVENTION: Colon Cancer Antigen Panel
FILE REFERENCE: L0461/7105 (JRV)
CURRENT APPLICATION NUMBER: US/09/849, 602
CURRENT FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: UNSURE
LOCATION: (76)..(76)
OTHER INFORMATION: X = any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCT-US02-21461-3; Sequence 3, Application PC/TUS0221461; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9, Application US/09860286
GENERAL INFORMATION:
APPLIENT: Gregory, Richard J.
                                                                                                           Sequence 24, Application US/09849602 GENERAL INFORMATION:
                                                                                                                                                          APPLICANT: Scanlan, Matthew J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSQETFSDLWKLLPEN 18
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    12 PPLSQETFSDLWKLLPEN 29
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Best Local Similarity 100.0
Matches 18; Conservative
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PCT-US02-21461-3
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Best Local Similarity
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US-09-849-602-24
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LENGTH: 393
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US-09-860-286-9
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ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: ASYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
FILING DATE: 18-May-2001
CLASSIFTCATION DATA:
PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

COMPATIBLE TO ADATA:

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Maneval, Daniel C.
TITLE OF INVENTION: Recombinant Adenoviral Vector and
Methods of Use
                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10274874
GENERAL INFORMATION:
APPLICANT: Trink, Barry
APPLICANT: Trink, Barry
APPLICANT: Saldransky, David
TITLE OF INVENTION: P40 Protein Acts as an Oncogene
FILE REFERENCE: 0100'7965
CURRENT APPLICATION NUMBER: US/10/274,874
CURRENT FILING DATE: 2002-10-22
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
                                                                                                                                                                                                                                                                                                                    STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1..393
OTHER INFORMATION: /note= "human p53"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION UNMERR: US/08/328,673
FILING DATE: 25-Oct-1994
APPLICATION NUMBER: US 08/142,669
FILING DATE: 25-OCT-1993
APPLICATION NUMBER: US 08/233,669
FILING DATE: 26-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Smith, Timothy S. REGISTRATION NUMBER: 35,367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 393 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: California
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Sequence 141, Application US/10211088
GENERAL INFORMATION:
APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
APPLICANT: Chen, Yih-Tai
TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding
CURRENT APPLICATION NUMBER: US/10/211,088
CURRENT FILING DATE: 2002-10-15
PRIOR PILLOATION NUMBER: 60/309,395
PRIOR APPLICATION NUMBER: 60/341,589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: Peptide
US-60-438-805-9
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US-10-276-774-2505
i Sequence 2505, Application US/10276774
j GENERAL INFORMATION:
i APPLICANT: Hyseq, Inc.
i APPLICANT: Hyseq, Inc.
i TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
j TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
j FILE REFERENCE: 21272-030
j CURRENT APPLICATION NUMBER: US/10/276,774
cURRENT FILING DATE: 2000-11-18
j PRIOR PILING DATE: 2000-04-27
pRIOR FILING DATE: 2000-04-27
pRIOR PLING DATE: 2000-02-03
j NUMBER OF SEQ ID NOS: 2700
software: Custom
j SOFTWARE: Custom
j SOFTWARE: Custom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 86.7%; Score 91; DB 6; Length 32; 94.4%; Pred. No. 1.1e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                87.6%; Score 92; DB 7; Le
100.0%; Pred. No. 4.5e-08;
tive 0; Mismatches 0;
                                         Sequence 9, Application US/60438805
GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS
TITLE OF INVENTION: MIXED ELEMENT CAPTURE AGENTS
FILE REFERENCE: UTSD:995USP1
CURRENT APPLICATION NUMBER: US/60/438,805
CURRENT APPLICATION NUMBER: 2003-01-09
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-276-774-2505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-211-088-141
                           US-60-438-805-9
                                                                                                                                                                                                                                              SEQ ID NO 9
LENGTH: 19
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING MAJOR DEPRES
TITLE OF INVENTION: DISORDER

TITLE OF INVENTION: DISORDER

TITLE OF INVENTION: DISORDER

TITLE OF INVENTION: DISORDER

CURRENT APPLICATION NUMBER: US/60/423,552

CURRENT FILING DATE: 2002-11-05

NUMBER OF SEQ ID NOS: 256

SEQ ID NO 118

LENGTH: 393
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING MAJOR DEPREG TITLE OF INVENTION: DISORDER.

TITLE OF INVENTION: DISORDER.

TITLE OF INVENTION: DISORDER.

CURRENT APPLICATION NUMBER: US/60/427,579

CURRENT FILING DATE: 2002-11-20

NUMBER OF SEQ ID NOS: 256

SOFTHARE: Patentin version 3.1

SEQ ID NO 118

LENGTH: 393
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.3%; Score 99; DB 7; Length 393; 100.0%; Pred. No. 7e-08; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%; Score 99; DB 7; Length 393; 100.0%; Pred. No. 7e-08; Live 0; Mismatches 0; Indels
PRIOR FILING DATE: 1998-03-27
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 7
US-60-423-552-118
; Sequence 118, Application US/60423552
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-60-427-579-118; Sequence 118, Application US/60427579; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                              1 PPLSQETFSDLWKLLPEN 18
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Best Local Similarity 100.
Matches 18; Conservative
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                                                                                   ; SEQ ID NO 4
; LENGTH: 393
; TYPE: RT
; CRGANISM: Homo sapiens
US-10-274-874-4
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; ORGANISM: Homo sapiens
US-60-423-552-118
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Matches
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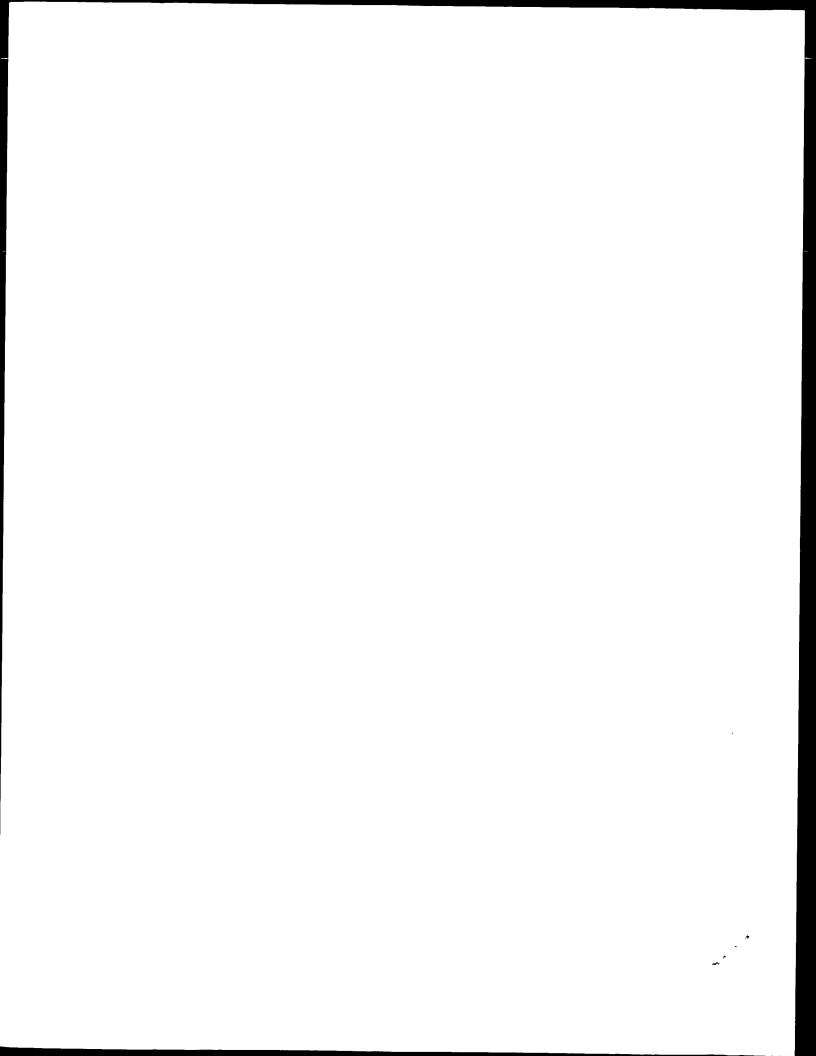
Gaps

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; OTHER INFORMATION: Description of Artificial Sequence:derived from p53
US-09-701-080C-27
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APPLICANT: Leach, Martin D.
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: NOVEL POLYMUCLECTIDES AND POLYPEPTIDES
TITLE OF INVENTION: ENCORED THEREBY
FILE REFERENCE: 1596-543 CON
CURRENT APPLICATION NUMBER: 09/540,763
PRIOR APPLICATION NUMBER: 09/540,763
PRIOR PILING DATE: 2000-03-30
PRIOR PILING DATE: 1999-04-05
PRIOR PILING DATE: 1999-04-05
PRIOR FILING DATE: 1999-04-02
PRIOR FILING DATE: 1999-03-11
PRIOR FILING DATE: 1999-03-11
NUMBER OF SEQ ID NOS: 6322
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Southwood, Scott
APPLICANT: Epimmune, Inc.
1TILE OF INVENTION: HLA Binding Peptides and Their Uses
ITILE REFERENCE: 39963-20019, 20
CURRENT FILING DATE: 1998-111-10
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 286
LENGTH: 11
                                                                                                    Length 13;
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                                                                                                  Query Match 66.7%; Score 70; DB 5; Length 13; Best Local Similarity 100.0%; Pred. No. 9.6e-05; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: p53.14 peptide 34.0324
US-09-189-702A-286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 250, Application US/10218140; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
                                                                                                                                                                                                                                       1 SQETFSDLWKLLP 13
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US-09-189-702A-286
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US-10-218-140-250
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LENGTH: 927
     FEATURE:
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US-09-701-080C-27
Sequence 27, Application US/09701080C
Sequence 27, Application US/09701080C
GENERAL INFORMATION
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE SPERENCE: N73477C GGW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: GB 9811303.8
PRIOR FILING DATE: 1998-05-26
PRIOR FILING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22, Application US/09701080C
GENERAL INFORMATION:
APPLICANT: INSTITUTE
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 J
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REFERENCE: N7347C GW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR APPLICATION NUMBER: GB 990157.0
PRIOR PLING DATE: 1998-05-26
PRIOR PLING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN VEY: 2.1
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                                                                                                                                                                                                 ; OTHER INFORMATION: Binding domain US-10-211-088-141
PRIOR FILING DATE: 2001-12-13
NUMBER OF SEQ ID NOS: 366
SOFTWARE: Patentin version 3.1
SEQ ID NO 141
                                                                                                                                                        ORGANISM: Artificial sequence
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Query Match 49.0%; Score 51.5; DB 6; Length 927;
Best Local Similarity 47.4%; Pred. No. 6;
Matches 9; Conservative 6; Mismatches 3; Indels 1; Gaps
Qy 2 PLSQB-TFSDLWKLLPENG 19
Db 616 PLEQDSTFAELWRTISKNG 634
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Search completed: February 12, 2003, 10:42:49 Job time : 20 secs



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Sequence 8, Appli
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                                                                  February 12, 2003, 10:35:42; Search time 11 Seconds (without alignments) 44.130 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/USOB_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
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11: /cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
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14: /cgn2_6/ptodata/1/pubpaa/USO0_NEW_PUB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-10-155-059-8
US-10-155-059-10
US-10-155-059-11
US-10-155-059-11
US-09-732-344-3
US-09-732-344-3
US-09-214-371-73
US-09-214-371-73
US-09-214-371-73
US-09-214-371-73
US-09-214-371-73
US-09-214-371-17
US-10-155-059-19
US-09-214-371-1
                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                 140259 seqs, 25548876 residues
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Maximum Match 100%
Listing first 45 summaries
                                             OM protein - protein search, using sw model
                                                                                                                                      1 PPLSQETFSDLWKLLPENG 19
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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Match Length DB
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Sequence 130, App
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Sequence 7, Appli
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Sequence 20, Appl
Sequence 21, Appl
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Sequence 24, Appl
Sequence 8, Appli
                                                                                                Sequence 21, App.
                                                                          Sequence 172, App
Sequence 12, Appl
                                               Sequence 26, 1
Sequence 33, 1
Sequence 34, 1
Sequence 17, 1
                  Sequence 39,
                              Sequence
                                       Sequence
          Sequence
              Sequence
      Sequence
                         US-09-214-371-7

US-09-214-371-24

US-09-214-371-41

US-09-214-371-25

US-09-214-371-25

US-09-214-371-33

US-09-214-371-33

US-09-019-679-3

US-09-019-679-3

US-10-078-770-172
US-10-155-059-18
US-09-214-371-17
US-09-950-692-6
US-10-155-059-22
US-09-214-371-39
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US-09-214-371-20
US-09-905-129-21
US-09-991-630-21
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US-10-155-059-24
                     US-10-024-123-7
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# ALIGNMENTS

Db 1 PPLSQETFSDLWKLLPEN 18 RESULT 2

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78210-12 7.8210-155-059-8 7.8equence 8, Application US/10155059 7.Patent No. US20020147173A1

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TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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             COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION OF CURNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
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COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIPICATION: <UNknown>
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APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-WAY-1998
PAPLICATION NUMBER: 60/046,207
FILING DATE: 12-WAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                   NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFRENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPRAS: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; SEQUENCE DESCRIPTION: SEQ ID NO: 10: US-10-155-059-10
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 amino acids
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                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
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Jost, Christine
TITLE OF INVENTION: METHODISS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                            TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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94.3%; Score 99; DB 12; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.2e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FESSESO for Windows Version 2.0b
SOFTWARE: PESSESO for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-AMY-2002
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRION APPLICATION DATE: US/09/081,975
PILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Eisenstein, Ronald I
REGISTRATION UNDERS: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                           ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 617-345-1300
     GENERAL INFORMATION:
APPLICANT: Kaelin, William
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 8:
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                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: MA
COUNTRY: USA
ZIP: 02110
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MUTATIONS WHICH SUPPRESS P53 CANCER MUTA- TIONS
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         94.3%; Score 99; DB 12; Length 20; 100.0%; Pred. No. 6.2e-09; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/776,695
FILING DATE: 26-Feb-2001
CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                    Sequence 32, Application US/09776695; Patent No. US20020068283A1; CENERAL INFORMATION:
APPLICANT: Boeke, Jef
Brachmann, Rainer
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING P53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILB REPERENCE: 21508-044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 01107.03170 TELECOMMUNICATION INFORMATION:
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US-09-776-695-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
PAPLICATION NUMBER: 08/795,006
FLING DATE: 2001-03-28
ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Banner & Witcoff STREET: 1001 G Street, NW CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 393 amino acids
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INFORMATION FOR SEQ ID NO: 32:
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                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 32
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                                                                                                                                    2 PPLSQETFSDLWKLLPEN 19
                                                                                                    1 PPLSQETFSDLWKLLPEN 18
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                                Best Local Similarity 100.
Matches 18; Conservative
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Best Local Similarity 100.
Matches 18; Conservative
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US-09-776-695-32
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           Query Match
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Jost, Christine
Jost, METHONS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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94.3%; Score 99; DB 12; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.2e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <u >Unknown></u>
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APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                   TYPE: amino acid
STRANDENESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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TELEPHONE: 617-345-6054
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Patent No. US20020147173A1
GENERAL INFORMATION:
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COMPUTER: IBM Compatible
                                                                                                    INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 617-345-1300
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STRANDEDNESS: single
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INFORMATION FOR SEQ ID NO: 16:
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ZIP: 02110
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US-10-155-059-16
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APPLICANT: Gregory, Ren N.
Wills, Ken N.
Maneval, Daniel C.
TITLE OF INVENTION: Recombinant Adenoviral Vector and
Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: US/09/860,211
FILING DATE: 18-May-2001
CLASSIFICATION: <a href="Chicago: Chicago: Chicago:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
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TELECOWHUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: 1..393

DTHER INFORMATION: /note= "human p53"
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APPLICATION NUMBER: US 08/328,673

FILING DATE: 15-FEB-2000

APPLICATION NUMBER: US 08/142,669

FILING DATE: 25-OCT-1993

APPLICATION NUMBER: US 08/233,669

FILING DATE: 26-APR-1994

ATTORNEY/AGENT INFORMATION:

NAME: Smith, Timothy S.

REGISTRATION NUMBER: 35,367
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR PILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 393 amino acids
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STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/09860211
Patent No. US20020137212A1
GENERAL INFORMATION:
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 9:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 PPLSQETFSDLWKLLPEN 29
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Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                         393
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, Using
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: METHODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                         0
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                                                               94.3%; Score 99; DB 10; Length 393; 100.0%; Pred. No. 1.5e-07; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.3%; Score 99; DB 12; L. 100.0%; Pred. No. 1.5e-07; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMNUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFRAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Eisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-155-059-14 ; Sequence 14, Application US/10155059
                                                                                                                                                                                                                                                                                                        Sequence 4, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 394 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                       1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                 12 PPLSQETFSDLWKLLPEN 29
                                                                 Query Match
Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
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Matches 18; Conserv
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US-09-860-211-9
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APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 94; DB 12; Length 20;
Pred. No. 3.5e-08;
0; Mismatches 1; Indels
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                                       MEDIUM TYPE: Diskette
COMPUTER: IBM Comparible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 73, Application US/09214371B ; Patent No. US20010018511A1
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; OTHER INFORMATION: X = Biotin-Ser
US-09-214-371-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       89.5%;
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ORGANISM: Artificial Sequence
              COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PPLSQETFSDLWKLLPEN 18
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Best Local Similarity 94.47
Matches 17; Conservative
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LENGTH: 22
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                                            APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 15, Application US/10155059

Patent No. US20020147173A1

GENERAL INFORMATION:

APPLICANT: Kaelin, William

Jost, Christine

TITLE OF INVENTION: METHODS OF TREATMENT USING

NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE

ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASESEO for Windows
SOFTWARE: FASESEO for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION CURREN: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 310,628
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-10-155-059-14
                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                         NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPEN 18
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Patent No. US20020147173A1
                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 02110
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COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                        CITY: Boston
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Matches 17; Conserva
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Gaps

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ATTORNEY/AGENT INFORMATION:
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SEQ ID NO 1
                                                                                                                                                                                                                                                                                                  US-10-155-059-13
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                             Query Match
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                         0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 19;
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                         0; Indels
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SOFTWARE: FastSEQ for Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION - CURROWN>
PRIOR CLASSIFICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
. *FILING DATE: 12-May-1997
                                                                                                                                                                                                                                                            APPLICANT: Year, Zhi-Min
APPLICANT: Year, Zhi-Min
APPLICANT: Gu, Jijila
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR PILING DATE: 1999-12-08
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
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Best Local Similarity 100.0%; Pred. No. 6.6e-08;
Matches 17; Conservative 0; Mismatches 0;
      Pred. No. 5.5e-08;
                         0; Mismatches
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                    ; Sequence 7, Application US/09732384; Patent No. US20020132977A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette
      Best Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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                                                                                              6 PPLSQETFSDLWKLLPE 22
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                         17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17;
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US-09-732-384-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84.8%; Score 89; DB 10; Length 19; 94.1%; Pred. No. 1.9e-07; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                    Length 19;
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88.9%; Pred. No. 1.3e-07;
tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Gardia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interac
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1997-07-04
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEPAX: 617-345-1300
TELEX: cUnknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CIRRACTERISTICS:
LENGTH: 19 aming acids
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                                                                                                                                                                                                                                                                             TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lane, David
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                     STRANDEDNESS: single
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ORGANISM: Artificial Sequence
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SOFTWARE: Patentin Ver. 2.0
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Best Local Similarity 94.1'
Matches 16; Conservative
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Matches 16; Conservative
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February 12, 2003, 10:32:16; Search time 36 Seconds (without alignments) 70:327 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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2. SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:\*

3. SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:\*

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5. SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:\*

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7. SIDS2/gcgdata/geneseq/geneseqp-embl/AA198.DAT:\*

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13: SIDS2/gcgdata/geneseq/geneseqp-embl/AA199.DAT:\*

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Disclosure, Fig 1; 52pp; English.

-termi p53, p53 h	Human p53 fragment N-terminal region N-terminal region	uman MDM2	lian	Mammalian two-hybr	Mammalian two-hybr	p53 an	dia Sign	קי כנק	1	10 Pt Ct	p53 prot	053	ric p53 prot	ied p53	fied p	fied p53	fied p53 vari	1ed p53	cled p53	a pos	4 F	אמאים ייסוקיים ה		היי היי	, 4	in.
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## ALIGNMENTS

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p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
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                                                                                  AAW82321 standard; Peptide; 19 AA.
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                                                                                                                                                                                                                                                                                            22-FEB-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         p53 homologue TIP peptide.
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RESULT 1
                                                   AAW8232
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X8888888888888X8

This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 reduction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

19 AA; Sequence

ö 0; Gaps 100.0%; Score 105; DB 19; Length 19; 100.0%; Pred. No. 5.7e-10; ive 0; Mismatches 0; Indels C Best Local Similarity 100.0 Matches 19; Conservative Query Match

1 PPLSQETFSDLWKLLPENG 19 1 PPLSQETFSDLWKLLPENG 19 ò 셤

RESULT 2

AAW82319 standard; Peptide; 19 AA. AAW82319

AAW82319;

22-FEB-1999 (first entry)

p53 homologue TIP peptide.

p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.

Synthetic.

W09847525-A1.

29-OCT-1998.

98WO-GB01144. 20-APR-1998;

97GB-0008092. 22-APR-1997;

(UYDU-) UNIV DUNDEE.

Lane DP;

WPI; 1998-609932/51. 

New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2

Disclosure, Fig 1; 52pp; English.

This sequence is a peptide homologue of a region of p53 which binds to made. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in apoulation of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not oversepses mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

19 AA; Sequence

Query Match

100.0%; Score 105; DB 19; Length 19;

1 PPLSQETFSDLWKLLPEN 18

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This represents a p53 N-terminal peptide fragment used in an Elisa TIP casay for analysing the interaction between human encogenic protein MDM2 assay for analysing the interaction between human encogenic protein MDM2 and p53. The invention provides peptide derivatives capable of binding to the human MDM2 These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated company of MDM2. The peptides may be used to identify molecules that bind levels of MDM2. The poptides may be used to identify and design inhibitors of MDM2, p53 binding. They to MDM2 and to identify binding partners especially MDM2, diagnose consistents and for treatment or prevention of disease involving p53/MDM2 interactions. Sepecially tumours and viral infections. The peptides can interactions and viral infections. The peptides can consider and all capability. Rectally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can accivate p53 function and commandiation in normal cells. The peptides which mind the MDM2 binding site in p53, have a significantly greater blocking activity compared contribute protein with wild-type p53.
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                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents
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Lane D, Picksley S;
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                            Indels
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           5.7e-10;
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           Pred. No.
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100.0%; Pic
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Best Local Similarity 100.0
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                              Conservative
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               Best Local Similarity
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                                      19;
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                                    Matches
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Antibodies specific for p53 are detected by binding to immobilised
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                                                                                                                                                                                                                                                                                          fragments of the p53 gene product containing the antibody-binding region. Preferred fragments contain amino acids 1-241, 40-349, 40-389, 66-241, 66-393, 237-349, 237-393 and esp. 9-33, 37-52 or 368-386. See AAR51872-R51881 for sequences of these fragments.
                                                                                                                                                                                      Non-radioactive detection of p53 specific antibodies - by capture on immobilised p53 or its fragments, then reaction with labelled second antibody, for diagnosis of tumours and suitable for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; p53; antibody; detection; biosensor; cancer patient;
p53 gene therapy; immune response; mutant.
                                                                                                                               Zentgraf H;
                                                                                                                                                                                                                                                                                                                                                                                 94.3%; Score 99; DB 15; Length 25; 100.0%; Pred. No. 7.1e-09; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p53 peptide used to detect antibodies against p53
                                                                                                                            Volkmann M,
                                                                                                (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM STIFTUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW60202 standard; peptide; 26 AA.
                                                                                                                            Tessmer C,
                                                                                                                                                                                                                                                  Claim 11; Page 19; 35pp; German.
                                                  93WO-EP02666.
                                                                           92DE-4232823
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                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 PPLSQETFSDLWKLLPEN 21
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Best Local Similarity 100."
Matches 18; Conservative
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                                                                                                                           Klein R, Schranz P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SCHE ) SCHERING CORP
                                                                                                                                                WPI; 1994-135732/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1998-240965/21.
                                                                                                                                                             N-PSDB; AAQ62365.
                                                                                                                                                                                                                                                                                                                                                           25 AA;
                                                 30-SEP-1993;
                                                                          30-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-OCT-1997;
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                                                                                                                                                                                                                               screening
                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AAR54907-R54921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                                                                                 cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                        New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human nuclear phosphoprotein p53; tumour suppressor gene product; anti-oncogene; cancer; tumour; antibody binding region; epitope.
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                                                                                                                                     Immunodominant epitope from p53 N-terminal.
                                                               AAR54907 standard; peptide; 25 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR51879 standard; Protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                          Soussi T;
                                                                                                                                                                                                                                                                                                                        92FR-0013110.
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1 PPLSQETFSDLWKLLPEN 18
                                                                                                                (first entry)
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18s 18; Conservative
                                                                                                                                                                                                                                                                                                                                               EURO-) LAB EUROBIO SA
                                                                                                                                                                                                                                                                                                                                                                      Legros Y, Lubin R,
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1994-167463/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 AA;
                                                                                                                                                                                                                                                                                                                     02-NOV-1992;
                                                                                                                                                                                                                                                                                            02-NOV-1993;
                                                                                                               29-NOV-1994
                                                                                                                                                                                                                                           WO9410306-A.
                                                                                                                                                                                                                                                                    11-MAY-1994
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                                                                                       AAR54907;
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                                      RESULT 4
                                                   AARS4907
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Peptides AAW60202-05 are derived from human p53 protein. The present peptide corresponds to residues 11-35. The peptides are used in the method of the invention. Antibodies that bind to p53 protein are detected by immobilising a p53 peptide directly on to a flow cell of a sensor chip in a biosensor, treating the peptide with a sample of patient serum, diluted in buffer, and measuring binding of antibody to the peptide using the biosensor. The method is used to monitor cancer patients undergoing p53 gene therapy (to determine if an immune response has developed), and also to detect antibodies against mutant forms of p53.
                                                                                                                                                                    Score 99; DB 19; Length 26;
                                                                                                                                                                               100.0%; Pred. No. 7.4e-09; cive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                  AAB30523 standard; peptide; 32 AA.
Claim 4; Page 5; 41pp; English.
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                                                                                                                                                                        94.3%;
                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                  3 PPLSQETFSDLWKLLPEN 20
                                                                                                                                                                                                                                                                                                                                               06-MAR-2001 (first entry)
                                                                                                                                                                                   Sest Local Similarity 100.
Aatches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-006911/01.
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                                                                                                                                                  26 AA;
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The specification describes a method of synthesis of biologically active substances of determined structure directly in the cells of living organisms containing specific RNA or DNA sequence. The method is based on the hybridisation of two or more oligomers bound with biologically inactive substances to specific RNA or DNA in vivo in the cells of living organisms. After hybridisation of the oligomers, the biologically inactive precursors bound to the oligomers can interact with each to make the active form of the substances. This changing of properties is due to chemical reactions which bind the biologically inactive precursors through a chemical bond into a biologically active form of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel methods for the synthesis of biologically active compounds from inactive precursors in the cells of living organisms, useful for producing proteins or polynucleotides -
                                                         Biologically active compound; cellular metabolism; DNA replication; RNA transcription; RNA translation; RNA elongation; RNA processing; protein synthesis; protein processing; cellular differentiation; cell division; ion channel transmission; cellular protein; toxin; RNA transportation; cellular oxidation; plasminogen antigen activator.
                             Peptide fragment of a tumour suppressor protein p53.
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the whole compound. The methods are useful for producing biologically active compounds from inactive precursors. These compounds may be inhibitors or stimulators of callular metabolism, DNA replication, RNA translation, RNA elongation RNA processing, protein synthesis, protein processing, cellular differentiation, cell division, ion channel transmission, cellular protein and RNA transportation, processes of cellular oxidation, toxins, proteins or RNAs. AAB30523-36 processes of cellular oxidation, toxins, proteins or RNAs. AAB30523-36 processes of cellular oxidation, toxins, proteins or RNAs. AAB30523-36 processes of cellular oxidation, toxins, proteins or RNAs. AAB30523-36 represent peptides which are bound to oligomers AAC62167-80 and AAC62181-94. The peptides are fragments of the tumour suppressor protein p53 and invention is used to produce the tumour suppressor protein p53 from the bound peptides and oligomers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse, Mdm2, murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; scomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 wild type human p53 protein. This peptide corresponds to amino acids wild type human p53 protein. This peptide corresponds to amino acids 6-41 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murino double minute-2 (mdm2) protein (AAM13600). Mdm2 protein is a 90 KD phosphoprotein which binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein itself has oncogenic properties, especially in a p53-mull background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The peptides AAW13602-6 represent peptide fragments derived from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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100.0%; Pred. No. 9.2e-09;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p53 protein amino acids 6-41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32 AA;
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constructed from the wild type human p53 sequence.

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over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukaemia; Ecell lymphoma, or other hyperpoliferative conditions such as restenosis. Note: this sequence is not given in the specification but is constructed from the wild type human p53 sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 36;
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hes 0; Indels
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Pred. No. 1.1e-08;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.3%; S
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RHON ) RHONE POULENC RORER SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dubs-Poterszman M, Tocque B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4; Page -; 43pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            p53 protein amino acids 1-41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95FR-0010331.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7 PPLSOETFSDLWKLLPEN 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JAN-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   contg. this nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-192837/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                    36 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9709343-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW13603;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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The peptides AAW13602-6 represent peptide fragments derived from the wild type human p53 protein. This peptide corresponds to amino acids 1-52 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double composed to bind the N-terminal amino acids (1-134) of the murine double composed to bind the N-terminal amino acids (1-134) of the murine double composed to bind the N-terminal amino acids (1-134) of the murine double composed to bind the N-terminal amino acids (1-134) of the murine double composed to binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdma protein citself has oncogenic properties, especially in a p53-null background. Composed to unblock cell cycle arrest in G1 caused by cover-expression of the p107 protein. The p53 spetides are examples of antagonists of the invention which are able to inhibit the oncogenic contivity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukaemia; B coll lymphoma, or other hyperproliferative conditions such as constructed from the wild type human p53 sequence.
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                                                                                                                                                                                                                                                                                                                                                                               tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                                                                                                                                                                                                                                                                           Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation;
                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating cancer with antagonist of oncogenic activity of protein Mdm2 - or nucleic acid encoding an antagonist, also viral vectors contg. this nucleic acid
                                       ;
0
   Length 41;
                                     0; Indels
Score 99; DB 18; I
Pred. No. 1.2e-08;
                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INRM ) INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                         AAW13602 standard; peptide; 52 AA.
               100.0%; Pr
tive 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4; Page -; 43pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dubs-Poterszman M, Tocque B,
                                                                                                                                                                                                                                                                                                                         p53 protein amino acids 1-52.
94.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96WO-FR01340.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95FR-0010331.
                                                                     1 PPLSOETFSDLWKLLPEN 18
                                                                                               12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                   16-JAN-1998 (first entry)
                                 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-192837/17.
               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      52 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9709343-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-SEP-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                           restenosis.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                 Matches
                                                                                                                                                                     RESULT 10
                                                                                                                                                                                     AAW13602
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94.3%; Score 99; DB 18; Length 52; 100.0%; Pred. No. 1.6e-08;

Best Local Similarity

Query Match

us-09-403-440a-2.open.rag

AAW07886;

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                                                                                                                                                                                                                                                      p53 gene; tumour suppressor gene; regulation; cellular proliferation; cellular transformation; carcinoma; human; tumour; MDM2; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosing neoplasia from amplification of MDM2 gene - or elevated gene expression, also new DNA, MDM2 protein, antibodies and treatment of sarcoma by inhibiting MDM2 expression
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%; Score 99; DB 14; Length 64; 100.0%; Pred. No. 2e-08; ive 0; Mismatches 0; Indels
0; Indels
                                                                                                                                                                                                                                                                                                                                                                  13..41
/note= "Site of interaction with MDM-2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kinzler KW, Vogelstein B;
 Mismatches
                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 35; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW07886 standard; protein; 64 AA.
                                                                                                                                       AAR42174 standard; peptide; 64 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93WO-US03199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92US-0867840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     92US-0903103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYJO ) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 PPLSQETFSDLWKLLPEN 29
                              1 PPLSQETFSDLWKLLPEN 18
                                                 12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                 05-MAY-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18; Conservative
    Conservative
                                                                                                                                                                                                                              p53 N-terminal fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Burrell M, Hill DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1993-336944/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                          gene amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                    Binding-site
                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       23-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                              WO9320238-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                           14-OCT-1993.
    18;
                                                                                                                                                                      AAR42174;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 12
AAW07886
ID AAW078
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    Matches
                                                                                                          RESULT 11
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AAW07886 represents the human p53 protein which is involved in the development of many cancers. The protein is used here in a method for identifying compounds that interfere with the binding of p53 and MDM-2. In binding the p53 protein, the MDM-2 protein releases a cell from p53-regulated growth, allowing cancers to develop. Therefore compounds identified as interfering with the binding of MDM-2 to p53 are potentially useful in the treatment of human neoplastic cells. In the method pref. one or both of the proteins is a fusion protein esp. with an antibody or antibody fragment which aids separation and identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identification of cpds. interfering with human MDMZ/p53 binding useful as therapeutic agents to treat human neoplastic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
                                                                                          p53; MDM-2; binding-inhibitor; identification; tumour; cancer; neoplasia; antibody fusion protein; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%; Score 99; DB 17; Length 64;
100.0%; Pred. No. 2e-08;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                           13..57
'note= "MDM-2 binding fragment"
                                                                                                                                                                                                                            /note= "MDM-2 binding fragment"
                                                                                                                                                                                                  /note= "MDM-2 binding fragment"
                                                                Human p53, involved in tumour suppression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 16; Column 19-20; 36pp; English.
                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW57240 standard; protein; 64 AA.
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                                                                                                                                                                                                                                                                                                                                                                                       92US-0867840.
                                                                                                                                                                                                                                                                                                                                                                                                       92US-0903103.
                                                                                                                                                                                                                                                                                                                                              92US-0867840.
                                                                                                                                                                                                                                                                                                                                                                           93US-0044619
                                                                                                                                                                                                                                                                                                                                                                                                                   94US-0245500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYJO ) UNIV JOHNS HOPKINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kinzler KW, Vogelstein B;
                                         28-JAN-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-401591/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
nes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    64 AA;
                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                               07-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                      18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                       07-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                        23-JUN-1992;
                                                                                                                                                                                                                                                                                        US5550023-A.
                                                                                                                                                                                                                                                                                                                    27-AUG-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                       Region
                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                              Region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 13
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The present sequence represents a human p53 fragment 13-41 which is necessary for the interaction of MDM-2 and p53. The present invention describes a method for diagnosing a neoplastic disease caused by overexpression of MDM2 protein. The method comprises detecting an elevated cellular amount of this protein. The method is useful for the diagnosis of sarcoma, especially liposarcoma, malignant fibrous histiocytoma or osteosarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human MDM2 binding polypeptide - comprises fragments of p53, useful in re-establishing p53-regulated growth control in cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MDM2; tumour; diagnosis; neoplasia; DNA binding protein;
p53 polypeptide; binding; tumour cell; p53-regulated growth;
inhibition; anti-cancer agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 99; DB 19; Length 64;
Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                    Cancer diagnosis - by determination of MDM2 protein
                                                                                                                                                                                                                         Burrell M, Hill DE, Kinzler KW, Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-terminal region of p53 (amino acids 13-41).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                           Disclosure; Column 19-20; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.3%; Scur
100.0%; Pre
0; 1
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                                                        95US-0390517
                                                                                              93US-0044619
                                                                                                             92US-0867840.
                                                                                                                                                     95US-0390517
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                                                                                                                                                                                      SNINGO UNIV ( OLYU)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Burrell M, Hill DE,
                                                                                                                                                                                                                                                                WPI; 1998-239206/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 18; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-100408/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64 AA;
                                                        17-FEB-1995;
                                                                                                             07-APR-1992;
23-JUN-1992;
                                                                                                                                                     17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-1992;
23-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW42878;
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셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method has been developed for inhibiting the growth of tumour cells containing a human MDM2 gene amplification. The method comprises treating the tumour cells with a DNA molecule that expresses a crashing to tuman MDM2 protein. The present collypeptide capable of binding to human MDM2 protein. The present invention describes three preferred human MDM2 protein. The present invention describes three preferred polypeptides for binding human MDM2: (1) the polypeptide comprises amino acids 1-50 of p53 (see AAW77240); (2) the polypeptide comprises amino acids 1-41 of p53 (see AAW77240) and at least none additional p53 residues on the N- or C-terminal side, provided that the polypeptide comprises amino acids 13-41 of p53 (see AAW57241) and at least nine additional p53 residues on the N- or C-terminal side, provided that the polypeptide comprises amino acids 13-41 of p53 (see AAW57241) and at least nine additional p53 residues on the N- or C-terminal side, provided that the polypeptide lacks amino acids 138-193 of p53. Some malignant fibrous hieriocytomas and liposarcomas have an MDM2 gene amplification, so detection of increased expression of MDM2 gene products indicates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibiting growth of tumour cells having MDM2 gene amplification with p53 protein fragment
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       Human; p53; MDM2; tumour; growth inhibition; amplification;
malignant fibrous histiocytoma; liposarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94.3%; Score 99; DB 19; Length 64; 100.0%; Pred. No. 2e-08; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Column 19-20; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW48243 standard; peptide; 64 AA.
                                                                                                                                                                                                               93US-0044619.
92US-0867840.
92US-0903103.
95US-0390515.
                                                                                                                                                                              95US-0390515.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%;
                                                                                                                                                                                                                                                                                                          SNINGO NINU ( OCYU)
                                                                                                                                                                                                                                                                                                                                                     Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 PPLSQETFSDLWKLLPEN 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human p53 fragment 13-41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-321574/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     64 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tumourigenesis.
                                                                Homo sapiens
                                                                                                                                                                              17-FEB-1995;
                                                                                                  US5756455-A.
                                                                                                                                                                                                                 07-APR-1993;
                                                                                                                                                                                                                                                                                                                                               Kinzler KW,
                                                                                                                                                                                                                                     07-APR-1992;
                                                                                                                                                                                                                                                                        17-FEB-1995
                                                                                                                                         26-MAY-1998
                                                                                                                                                                                                                                                         23-JUN-1992
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Gaps

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us-09-403-440a-2.open.rag

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The present sequence represents a N-terminal portion of the p53 protein. These amino acid residues have been found to be necessary for the interaction of MDM2 and p53. A cell containing three recombinant but constructs was produced. These constructs encode an MDM2 protein fused to a sequence-specific DNA binding domain, a p53 polypeptide fused to a transcriptional activation domain, and a reporter gene downstream from a DNA element which is recognised by the sequence-specific DNA-binding domain. The cell is used to identify a compound which interferes with the binding of MDM2 and p53. Since MDM2 is overexpressed in tumour cells and since binding of MDM2 to p53 appears to allow tumour cells to escape from p53-regulated growth, compounds that inhibit such binding would be useful as anti-cancer agents.
                                                  Claim 1; Columns 19-20; 41pp; English.
  over-expressing MDM2
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 64 AA;
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Ouery Match

94.3%; Score 99; DB 19; Length 64;
Best Local Similarity 100.0%; Pred. No. 2e-08;
Matches 18; Conservative 0; Mismatches 0; Indels 1 PPLSOETFSDLWKLLPEN 18 12 PPLSQETFSDLWKLLPEN 29 g ઠે

0; Gaps

Search completed: February 12, 2003, 10:34:44 Job time : 37 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:33:46; Search time 15 Seconds (without alignments) 121.770 Million cell updates/sec Run on:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283224 seqs, 96134422 residues Searched:

Total number of hits satisfying chosen parameters:

283224

length: 0 length: 2000000000 Minimum DB seq Maximum DB seq

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR\_73:\*
1: pirl:\*
2: pir2:\*
3: pir3:\*
4: pir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

SUMMARIES	DB I	91 2 JC6193 tumor suppressor p	1 DNHU53 cellul	1 \$06594 cellular tumor	1 S51648 cellular tumor	2 JC6176	96 1 JH0633	63 1 A29376 cellular tumor	2 538824	1 DNMS53 cellular tumor	1 S02192	1 JH0631	2 \$28857	2 JH0266 glutamate	2 C64396 precorrin	2 G86715	2 E96672	2 C49921	2 AD3262	2 B25039	1 VCTMPV coat protein - p	2 G84779 hypothetical p	2 D83027 hypothetical	2 T41241 oxysterol-bin	2 G70327	0	2 C75460 hypothetical	2 S74792 hypothetical	
	Length	391	393	393	386	393	396	363	381	390	391	396	1009	1009	230	367	447	119	308	1004	157	469	487	489	926	250	485	274	COC
d	Query Match I	94.3	94.3	94.3	89.5	81.9	81.9	77.1	70.5	70.5	9.89	57.1	49.0	49.0	47.6	47.6	47.6	45.7	45.7	•	44.8	٠	٠	44.8	44.8	44.3	44.3	43.8	0 7
	Score	66	66	6	94	9	G		4	4	0	90		'n		0	20	48	48	48	47		47	47	4	46.5	46.5	46	,,
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E96796 AD2194	I57557 B95064	D97931 P3WLB4	503072	A82749 S61204	ASLJMS	S61205 S54852	S61206	S61208	S54853	S54849
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367	837	211	114	134 209	211	238 239	239	244	246	250
42.9	42.9 42.4	42.4	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9
45 45	45 44.5	44.5 44	4.	4 4 4 4	44	4 4 4 4	44	44	44	44
30 31	3 3 3 3	ა ი 4. ი	36	38	39	4.4. 0.1.	42	43	44	<b>4</b> 5

### ALIGNMENTS

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Cispecies: Oryctolagus cuniculus (domestic rabbit)
Cispecies: Nav. P.; Requence_revision 09-May-1997 #text_change 20-Jun-2000
Cispecies: Nav. P.; Romco, P.; de Fromentel, C.C.
Gene 185, 169-173, 1997
A;Title: CDNA cloning and immunological characterization of rabbit p53.
A;Reference number: JC6193; MUID:97208869; PMID:9055811
A;Reference number: JC6193; MUID:97208869; PMID:9055811
A;Residues: 1-391 <LEA>
A;Residues: 1-391 <LEA>
A;Cross-references: EMBL:X90592; NID:q1532043; PIDN:CAA62216.1; PID:q1532044
A;Genetica: p53
C;Superfamily: cellular tumor antigen p53
C;Keywords: tumor
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RESULT 1
                                         JC6193
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UNAULAR THORY ANTIGOR DESS [validated] - human
NyAlternate names: cellular phosphoprotein p53; oncoprotein p53; transformation suppress Cillular tumor antigen p53 [validated] - human
NyAlternate names: cellular phosphoprotein p53; oncoprotein p53; transformation suppress CiSpecies Home sapiens (man)
C;Species: Home sapiens (man)
C;Date: 05-0ct-1988 #sequence revision 18-Nov-1994 #text change 15-Sep-2000
C;Accession: A2524; A43073; JT0436; S40773; S42669; A22837; A55060; A25397; S42
A905; ISB346; IYB850; I52681; S60153
R;Lamb, P.; Crawford, L.
Mol. Cell. Biol. 6, 1379-1385, 1986
A;Title: Characterization of the human p53 gene.
A;Reference number: A25224; MUID:87064416; PMID:2946935
A;Residues: 1-33 < clambar
A;Residues: 1-333 < clambar
A;Residues: 1-333 < clambar
A;Reference number: JT0436; MUID:89108008; PMID:2905688
A;Title: A variation in the structure of the protein-coding region of the human p53 gene
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A;Residues: 1-333 < clambar
A;Residues: 1-333 < clambar
A;Residues: EMBL:M22898; NID:g189474

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A;Cross references: EMBL:X54156; NID:935213; PIDN:CAA38095.1; PID:935214
R;Yamada, Y.; Yoshida, T.; Hayashi, K.; Sekiya, T.; Yokota, J.; Hirohashi, S.; Nakatani,
Cancer Res. 51, 5800-5805, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Title: p53 gene mutations in gastric cancer metastases and in gastric cancer cell line
A,Reference number: A44905, MUID:92034678, PMID:1933850
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A)Residues: 1-253, 'D',255-393 <F115
A)Cross-references: EMBL:X60020; NID:9506452; PIDN:CAA42635.1; PID:9506453
A)Note: all sequences submitteed to the EMBL/GenBank/DDBJ databases June 1991
R)Futreal, P.A.; Barrett, J.C.; Wiseman, R.W.
Nucleic Acids Res. 19, 6977, 1991
A)Fitle: An Alu polymozphism intragenic to the TP53 gene.
A)Reference number: 138093; MUID:92107726; PMID:1762941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Residues: 1-245,'T',247-393 <F04>
A;Cross-references: EMBL:X60013; NID:g506438; PIDN:CAA42628.1; PID:g506439
A;Accession: 138086
A;Status: translated from GB/EMBL/DDBJ
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A;Cross-references: EMBL:X60014; NID:g506440; PIDN:CAA42629.1; PID:g506441
A;Accession: 138087
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A;Residues: 1-247,'Q',249-393 <F06>
A;Cross-references: EMBL:X60015; NID:9506442; PIDN:CAA42630.1; PID:9506443
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A;Cross-references: EMBL:X60016; NID:g506444; PIDN:CAA42631.1; PID:g506445
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A;Molecule type: mRNA
A;Residues: 1-247,'Q',249-393 <F08>
A;Cross-references: EMBL:X60017; NID:g506446; PIDN:CAA42632.1; PID:g506447
A;Accession: 138090
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A;Residues: 1-192,'R',194-393 <F02>
A;Cross-references: EMBL:X60011; NID:9506434; PIDN:CAA42626.1; PID:9506435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:X60012; NID:g506436; PIDN:CAA42627.1; PID:g506437
A;Accession: 138085
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    A;Cross-references: EMBL:X60010; NID:g506432; PIDN:CAA42625.1; PID:g506433
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A;Cross-references: EMBL:X60019; NID:g506450; PIDN:CAA42634.1; PID:g506451
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A,Residues: 246-247, W',249-250 <YAM>
A,Cross-references: GB:S63157; NID:g237829; PIDN:AAB20140.1; PID:g237830
A,Note: sequence extracted from NCBI backbone (NCBIN:63157, NCBIP:63158)
A,Note: mutation from a liver metastasis of a gastric cancer
R;Hensel, C.H.; Xiang, R.H.; Sakaguchi, A.Y.; Naylor, S.L.
                                                  A; Note: deletion of a C nucleotide causes a frameshift at position 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: mRNA
A;Residues: 1-71,'P',73-162,'H',164-393 <F09>
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A;Molecule type: mRNA
A;Residues: 1-393 <F03>
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A,Status: translated from GB/EMBL/DDBJ
A,Molecule type: mRNA
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A; Status: translated from GB/EMBL/DDBJ
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A;Status: translated from GB/EMBL/DDBJ
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A; Molecule type: DNA
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                                                                                       A;Accession: 138083
A;Status: translated from GB/EMBL/DDBJ
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A; Residues: 1-393 <CHU>
A; Cross-references: ENRI:X54156; NID:g35213; PIDN:CAA38095.1; PID:g35214
A; Cross-references: ENRI:X54156; NID:g35213; PIDN:CAA38095.1; PID:g35214
B; Matlashewski, G; Lamb, P.; Pim, D.; Peacock, J.; Crawford, L.; Benchimol, S.
ENBO J. 3, 3257-3362, 1984
A; Title: Isolation and characterization of a human p53 cDNA clone: expression of the hum A; Reference number: 842669; MUID:85126934; PMID:6396087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A Molecule type: mRNA

A; Residues: 1-71, Pr. 773-393 <ZAK>
A; Residues: 1-71, Pr. 773-393 <ZAK>
A; Residues: 1-71, Pr. 773-393 <ZAK>
A; Cross-references: ENBL:XO2469; EMBL:M60950; NID:G35209; PIDN:CAA26306.1; PID:G35210
R; Harlow, E.; Williamson, N.M.; Ralston, R.; Helfman, D.M.; Adams, T.E.
A) C. Cell. Biol. 5, 1601-1610, 1985
A; Title: Molecular cloning and in vitro expression of a cDNA clone for human cellular tu
A; Reference number: A55060; MUID:85267676; PMID:3894933
A;Note: this 72-Arg allele appears to be about 5 times more frequent than the 72-Pro all
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A;Residues: 1-189,'LLSILSEWKEICVWSIMMTETLFDIVWWCPMSRLRLALT','VPPSTTTTCVTVPAWAA' <F01>
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A; Cross-references: EMBL:X01405; NID:g35215; PIDN:CAA25652.1; PID:g642241
A; Cross-references: EMBL:X01405; NID:g35215; PIDN:CAA25652.1; PID:g642241
B; Zakut-Houri, R.; Bienz-Tadmor, B.; Givol, D.; Oren, M.
EMBO J. 4, 1251-1255, 1985
A; Title: Human p53 cellular tumor antigen: cDNA sequence and expression in COS cells.
A; Reference number: A22837; MUID:85230577; PMID:4006916
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A;Cross-references: GB:KOJ199; NID:g189479; PIDN:AAA59989.1; PID:g189479
A;Experimental source: Clone PR4-2, cell line A431
R;Harris, N.; Brill, E.; Shohat, O.; Prokocimer, M.; Wolf, D.; Arai, N.; Rotter, V. Mol. Cell. Biol. 6, 4650-4656, 1986
A;Title: Molecular basis for heterogeneity of the human p53 protein.
A;Reference number: A39086; MulD:87089926; PMID:3025664
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A; Residues: 1-71, P', 73-78, T', 80-393 < HAR2>
A; Residues: 1-71, P', 73-78, T', 80-393 < HAR2>
A; Residues: 1-71, P', 73-78
M2 Crosser-references: EMBL:M14695; NID:G339815
A; Experimental source: clone p53-H-19, transformed hybridoma SV-80 cell line
R; Matlashewski, G.J.; Tuck, S.; Pim, D.; Lamb, P.; Schneider, J.; Crawford, L.V.
Mol. Cell. Biol. 7, 961-963, 1987
A; Title: Primary structure polymorphism at amino acid residue 72 of human p53.
A; Reference number: S42452; MUID:87144273; PMID:3547088
                                                                                                                                                                      A,Cross-references: EMBL:M22898; NID:g189474; PIDN:AAAS9988.1; PID:g189476
A,Note: this 72-Pro allele was found in both normal and malignant cell lines
R;Chumakov, P.M.; Almazov, V.P.; Jenkins, J.R.
submitted to the EMBL Data Library, August 1990
A;Reference number: $40773
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A,Residues: 1-78, TT,80-393 <HARL>
A,Cross-references: EMBL:M14694; NID:g339813; PIDN:AAA61211.1; PID:g339814
A,Experimental source: clone p53-H-1, transformed hybridoma SV-80 cell line
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A;Residues: 66-71, 'P',73-79 <MKI2>
A;Experimental source: clone lambda C113
A;Note: 72-Cys was also found, and appears to represent a polymorphism
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A;Residues: 66-79 <MKI3>
A;Experimental source: clone J6K
Exferrell, P.J.; Allan, G.J.; Shanahan, F.; Vousden, K.H.; Crook, T.
EMBO J. 10, 2879-2887, 1991
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                                                                                            A;Molecule type: DNA
A;Residues: 1-71,'P',73-393 <BUC2>
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                                                       A; Accession: JT0436
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cellular tumor antigen p53 - bovine
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les 16; Conserva
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A; Residues: 1-393 <LEE>
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A;Residues: 1-386 <DEQ>
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A; Residues: 274-277, S',279-282 < HEN2>
A; Residues: 274-277, S',279-282 < HEN2>
A; Cross-references: GB $41977; NID:91679932; PIDN:AAB19325.1; PID:9232816
R; Chow, V. T.; Quek, H.H.; Tock, E.P.C.
Cancer Lett. 73, 141-148, 1993
A; Title: Alternative splicing of the p53 tumor suppressor gene in the Molt-4 T-lymphobla
A; Reference number: IS2681; MUID:94036762; PMID:8221626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; Molecule type: mRNA
A; Residues: 327-311, DQTSFQKENC' <CHO>
A; Cross-references: GB:S66666; NID:9436292; PIDN:AAB28601.1; PID:9436293
A; Note: mutant sequence with altered spliting and termination expressed in Molt-4 T-lymp
A; Note: mutant sequence with altered spliting and termination expressed in Molt-4 T-lymp
A; Note: mutant sequence with altered spliting and termination expressed in Molt-4 T-lymp
Mol. Gen. Genet. 249, 425-431, 1995
A; Title: Mapping of linear epitopes recognized by monoclonal antibodies with gene-fragme
A; Reference number: S60151; MUID:96133682; PMID:8552047
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A, Molecule type: mRNA

A, Residues: 1-393 «RIGA-

A, Residues: 1-393 «RIGA-

A, Cross-references: ENBL:X16384; NID:922795; PIDN:CAA34420.1; PID:922796

C, Superfamily: cellular tumor antigen p53

C, Superfamily: cellular tumor antigen p63

C, Superfamily: cellular tumor ontrol; DNA binding; homotetramer; nucleus; phosph

F, 176, 179, 238, 242/Binding site: zinc (Cys, Vsy, Cys) #status predicted

F; 392/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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R;Dang, C.V.; Lee, W.M.F.
J. Biol. Chem. 264, 18019-18023, 1989
A;Title: Nuclear and nucleolar targeting sequences of c-erb-A, c-myb, N-myc, p53, HSP70,
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C;Species: Cercopithecus aethiops (green monkey, grivet)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: 806594
R;Rigaudy, P.; Eckhart, W.
Nucleic Acids Res. 17, 8375, 1989
A;Title: Nucleotide sequence of a cDNA encoding the monkey cellular phosphoprotein p53.
                              A,Title: Use of the single strand conformation polymorphism technique and PCR to detect A,Reference number: 158354; MUID:91296386; PMID:1648702
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A;Residues: 244-247,'W',249-252 <HEN1>
A;Cross-references: GB:S41969; NID:g1679931; PIDN:AAB19324.1; PID:g232814
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A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                           A; Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local Similarity 100.
Matches 18; Conservative
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      Oncogene 8', 1067-1071, 1991
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Matches 18; Conserv
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                                                                                             Accession: 158354
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S51648

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A,Cross-references: EMBL:X81704; NID:g602332; PIDN:CAA57348.1; PID:g602333
C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; phosphoprotein; F;168,171,231,235/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;385/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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NiAlternate names: tumor-suppressor protein p53
C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: S51648
R;Dequiedt, F.; Willems, L.; Burny, A.; Kettmann, R.
Submitted to the EMBL Data Library, September 1994
A;Description: Nucleotide sequence of the ovine p53 tumor-suppressor gene cDNA and its (A;Reference number: S51648
A;Accession: S51648
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N;Alternate names: tumor-suppressor protein p53
C;Species: Mesocricetus auratus (golden hamster)
C;Dapeies: Mesocricetus auratus (golden hamster)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: UH0633
R;Legros, Y; Montyre, P.; Soussi, T.
A;Reference number: Discourage and immunological characterization of hamster p53.
A;Reference number: UH0633; MUID:92210007; PMID:1555773
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-396 <LEG>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 89.5%; Score 94; DB 1; Length 386; Best Local Similarity 94.4%; Pred. No. 1.7e-06; Matches 17; Conservative 0; Mismatches 1; Indels
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Pred. No. 2.8e-05;
0; Mismatches 1;
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C,Superfamily: cellular tumor antigen p53
C,Keywords: liver; tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            81.9%;
94.1%;
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A; Residues: 1-381 (ARA)
A; Residues: 1-381 (ARA)
A; Cross-references: GB:M13874; NID:g200202; PIDN:AAA39883.1; PID:g200203
R; Han, K.A.; Kulesz-Martin, M.F.
Nucleic Acids Res. 20, 1979-1981, 1992
Nucleic Acids Res. 20, 1979-1981, 1992
A; Title: Alternatively spliced p53 RNA in transformed and normal cells of different tiss A; Reference number: 835478; MUID:92253421; PMID:1579500
                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:M13874; NID:g200202; PIDN:AAA39883.1; PID:g200203
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, July 1388
C;Comment: This sequence, produced by alternative splicing of the tenth intron, lacks th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Residues: 1-134,'V',136-390 <BIE>
A;Cross-references: GB:X00876; NID:G871420; PIDN:CAA25420.1; PID:G871421; GB:X01237; GB:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    F;319-357/Region: tetramer association
F;79,12,12,18,23,37Binding site: phosphate (Ser) (covalent) #status predicted
F;173,176,235,239/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;312/Binding site: phosphate (Ser) (covalent) (by cdc2 kinase) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N;Alternate names: oncoprotein p53
C;Species: Mus musculus (house mouse)
C;Dete: 28-Mug-1985 #sequence_revision 04-Oct-1996 #text_change 11-May-2000
C;Date: 28-Mug-1985 #sequence_revision 04-Oct-1996 #text_change 11-May-2000
C;Accession: A22739, S06336; A02684, S38822; S38823; S40014; I48703
R;Bienz, B.; Zakut-Houri, R.; Givol, D.; Oren, M.
EMBO J. 3, 2179-2183, 1994
A;Title: Analysis of the gene coding for the murine cellular tumour antigen p53.
A;Reference number: A22739; MUID:85027173; PMID:6092064
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A;Title: Primary structure of DNA complementary to murine oncoprotein p53 mRNA.
A;Reference number: S06336; MUID:88221682; PMID:3329909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Molecule type: mRNA
A;Residues: 1-134, V',136-390 <CHU>
R;Zakut-Houri, R.; Oren, M.; Bienz, B.; Lavie, V.; Hazum, S.; Givol, D.
Nature 306, 594-597, 1983
A;Title: A single gene and a pseudogene for the cellular tumour antigen p53.
A;Reference number: A02684; MUID:84068204; PMID:6646235
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S'not known.
C'Superfamily: cellular tumor antigen p53
C'Keywords alternative splicing; phosphoprotein; zinc
C'Keywords alternative splicing; phosphoprotein; zinc
C'H-44/Domain: transcription activation #status predicted <TRA>
F;16-26/Region: conserved region I
F;16-12/Region: DNA-binding core #status predicted <DBC>
F;1108-121/Region: L1 loop
F;1108-121/Region: conserved region II
F;160-192/Region: conserved region II
F;121-252/Region: conserved region IV
F;231-252/Region: conserved region IV
F;231-252/Region: conserved region V
                                                                                                                                                                                                                                                                                                                                              nucleic acid sequence not shown; translation not shown
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A;Residues: 1-159,'H',161-167,'G',169-233,'I',235-390 <ZAK>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: not compared with conceptual translation
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                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Residues: 1-381 <HAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                            A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
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A; Accession: $38824
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cellular tumor antigen p53 - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Date: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession A29376; S61531; $72313; 151639
R;Soussi, T.; de Fromentel, C.C.; Mechali, M.; May, P.; Kress, M.
Oncogene 1, 71-78, 1987
A;Title: Cloning and characterization of a CDNA from Xenopus laevis coding for a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arresidues: 1-363 <SOU>
Arresidues: 1-363 <SOU>
Arross-references: EMBL:X05191; NID:g64961; PIDN:CAA28821.1; PID:g64962
Arross-references: EMBL:X05191; NID:g64961; PIDN:CAA28821.1; PID:g64962
Arrespons 9, 109-120, 1994
Arritle: Overexpression of wild-type p53 interferes with normal development in Xenopus 1
Arreference number: 151639; MUID:94134403; PMID:8302570
                                                                                                                                         C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph
F;179,182,241,245/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;395/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph
F;150,153,213,217/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;362/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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Cispecies: Mus musculus (house mouse)
Cispecies: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 23-Jul-1999
Cistocession: 838824; 835478
Fishari, N'; Nomura, D:; Yokota, K:; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V.
Mol. Cell. Biol. 6, 3232-3239, 1986
A;Title: Immunologically distinct p53 molecules generated by alternative splicing.
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A;Residues: 1-293,295-363 <HOE>
A;Residues: 1-293,295-363 <HOE>
A;Cross-references: EMBL:X77546; NID:g468513; PIDN:CAA54672.1; PID:g468514
R;Hoever, M.; Clement, J.; Wedlich, D.; Montenarh, M.; Knochel, W.
submitted to the EMBL Data Library, March 1994
A;Reference number: S72313
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A,Residues: 1-51,'S',53-70,72-293,295-363 <HOW>
A,Cross-references: EMBL:X77546; NID:g468513; PIDN:CAA54672.1; PID:g468514
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A;Cross-references: GB:M75144; NID:g191414; PIDN:AAA37085.1; PID:g191415
A;Experimental source: kidney, strain MP1
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                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 94.1%;
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Matches 14; Conservative
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                                                                                                                A;Gene:
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A;Cross-references: EMBL:X13058; NID:956828; PIDN:CAA31457.1; PID:956829
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A;Cross-references: EMBL:Z17238
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Best Local Similarity
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A;Status: translated from GB/EMBL/DDBJ
A;Residues: 1-47, 'R', 49-78, 'QW', 82-390 <RES>
A;Residues: 1-47, 'R', 49-78, 'QW', 82-390 <RES>
A;Cross-references: EMBL:X00741; NID:953570; PIDN:CAA25323.1; PID:953571
C;Comment: This DNA-binding protein plays an essential role in the regulation of cell di C;Comment: The tetramer association region may exhibit a beta-turn, beta-sheet, beta-tur C;Superfamily: cellular tumor antigen p53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: mRNA
A; Readdues: 1-167, (47, 169-390 <ARA3>
A; Cross-references: EMBL:M13873; NID:9200200; FIDN:AAA39882.1; PID:9200201
A; Cross-references: EMBL:M3873; NID:9200200; FIDN:AAA39882.1; PID:9200201
R; Jenkins, J.R.; Rudge, K.; Redmond, S.; Wade-Evans, A.
Nicleia Acids Res. 12, 5609-55626, 1984
A; Title: Cloning and expression analysis of full length mouse cDNA sequences encoding
A; Reference number: 148703; MUID:84272240; PMID:6379601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     F;319-357/Region: tetramer association
F;319-357/Region: tetramer association
F;7,9,12,18,23,37/Binding site: phosphate (Ser) (covalent) #status predicted
F;173,176,235,239/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;312/Binding site: phosphate (Ser) (covalent) (by cdc2 kinase) #status predicted
F;389/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
           A,Cross-xeferences: GB:X01237; GB:K01700; NID:g53575
R;Arai, N.; Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V.
Mol. Cell. Biol. 6, 323-3239, 1986
A;Title: Immunologically distinct p53 molecules generated by alternative splicing. A;Reference number: S38822; MUID:87064640; PMID:3023970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NiAlternate names: gene p53 protein; nuclear oncoprotein p53
C;Species: Rattus norregicus (Norway rat)
C;Species: Rattus norregicus (Norway rat)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: S02192; S41149
R;Soussi, T.; de Fromentel, C.C.; Breugnot, C.; May, E.
Nucleic Asids Res. 16, 11384, 1988
A;Title: Nucleotide sequence of a cDNA encoding the rat p53 nuclear oncoprotein.
A;Reference number: S02192; MUID: 89083585; PMID: 3060862
                                                                                                                                                                                                                                                                                                                                                                   A; Modecule type: mRNA
A; Residues: 1-167, 'G', 169-233,'I', 235-390 <ARA2>
A; Cross-references: BMBL:MI3873
R; Arai, N.; Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V.
submitted to the EMBL Data Library, July 1988
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A;Cross-references: EMBL:M13872; NID:g200198; PIDN:AAA39881.1; PID:g200199
A;Accession: S38823
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Fi99-289/Domain: DNA-binding core #status predicted <DBC>
Fi108-121/Region: L1 loop
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F;168-178/Region: conserved region III
F;231-252/Region: conserved region IV
F;233-248(Region: L3 loop
F;267-283/Region: conserved region V
F;313-319/Region: nuclear location signal
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A; Accession: S40014
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A, Molecule type: mRNA
A, Residues: 1-391 <SOU>
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Glutamate receptor delta-1 chain precursor - rat
NiAlternate names: Asinate receptor
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 20-Aug-1999
C;Accession: S28857; S31222
C;Accession: S28857; S31222
Fillowed, H.; Sprengel, R.; Laurie, D.J.; Koehr, G.; Herb, A.; Seeburg, P.H.; Wisden, W.FEBS Lett. 315, 318-322, 1993
A;Title: The rat delta-1 and delta-2 subunits extend the excitatory amino acid receptor
A;Reference number: S28857; MUID:93138096; PMID:8422924
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A.Introns: 25/2; 32/3; 123/3; 185/1; 259/2; 305/1; 329/3; 365/2
C;Superfamily: cellular tumor antigen p53
C;Reywords apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph
F;174,177,236,240/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;390/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-173, W',175-391 <HUL>
A;Residues: 1-173, W',175-391 <HUL>
A;Cross-references: EMBL:L07909
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, December 1992
C;Genetics:
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A,Status: preliminary, nucleic acid sequence not shown; translation not shown
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A; Title: Rainbow trout p53: cDNA cloning and biochemical characterization. A; Reference number: JH0631; MUID:92210006; PMID:1339362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cellular tumor antigen p53 - rainbow trout
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Date: 10-Sep_1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   68.6%; Score 72; DB 1; Length 391; 93.3%; Pred. No. 0.0037; tive 0; Mismatches 1; Indels
RiHulla, J.E.; Schneider, R.P.
Nucleic Acids Res. 21, 713-717, 1993
A;Title: Structure of the rat p53 tumor suppressor gene
A;Reference number: S41149; MUID:93181268; PMID:8441680
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0; Gaps

7; Indels

Length 230;

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Science 273, 1058-1073, 1996
A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.: Woese, C.A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii A;Reference number: A64300; MUID:96337999; PMID:8688087
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R;Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, K.; Weissenbach, J.; Ehrli
Genome Res. 11, 731-753, 2001
A;Title: The complete genome sequence of the lactic acid bacterium Lactococcus lactis ss
A;Reference number: A86625; MUID:21235186; PMID:11337471
                                                                                                                                                                                                                                                                                                                                              A;Residues: 1-230 <BUL>
A;Cross-references: GB:U67522; GB:L77117; NID:g2826315; PIDN:AAB98764.1; PID:g.592306;
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A;Cross-references: GB:AE005176; PID:g12723640; PIDN:AAK04825.1; GSPDB:GN00146
A;Experimental source: strain IL1403
                                                                                                                                                                                                            A, Accession: C64396
A, Status: preliminary; nucleic acid sequence not shown; translation not shown
A, Molecule type: DNA
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C;Date: 23-Mar_2001 #sequence_revision 23-Mar-2001 #text_change 03-Aug-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    racemase [imported] - Lactococcus lactis subsp. lactis (strain IL1403)
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55.6%; Pred. No. 4.5;
tive 1; Mismatches
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3; Mismatches
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C;Superfamily: precorrin-3 methylase
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Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                (Superfamily: glutamate receptor; glutamate receptor homology
(Keywords: glycoprotein; ion channel; neurotransmitter receptor; transmembrane protein
(F) 1-15/Domain: signal sequence #status predicted <SIG>
(F) 16-1009/Product: glutamate receptor delta-1 chain #status predicted <MAT>
(F) 16-1009/Product: glutamate receptor homology <GRHs
(F) 16-1009/Product: glutamate receptor homology <GRHs
(F) 16-1009/Product: glutamate receptor homology <GRHs
(F) 16-1009/Product: glutamate #status predicted <TM1>
(F) 16-100main: transmembrane #status predicted <TM3>
(F) 16-100main: transmembrane #status predicted <TM3>
(F) 16-100main: transmembrane #status predicted <TM3>
(F) 16-100main: transmembrane #status predicted <TM4>
(F) 16-100main: transmembrane 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 11-1009 <7AAA.
A;Cross-references: DDBJ:D1017; NID:g220417; PIDN:BAA01041.1; PID:g220418
A;Cross-references: DDBJ:D10171; NID:g220417; PIDN:BAA01041.1; PID:g220418
C;Comment: Glutamate receptor channels mediate most of the fast excitatory synaptic tranc; Superfamily: glutamate receptor; glutamate receptor; phosphoprotein; transmembrane prote [C;Keywords: glycoprotein; neurotransmitter receptor; phosphoprotein; transmembrane prote [F;1-15,Domain: signal sequence #status predicted <SIG>F;16-1009/Product: glutamate receptor channel delta-1 chain #status predicted <GLU>F;65-5-54/Domain: transmembrane #status predicted <TRL>F;65-54/Domain: transmembrane #status predicted <TRL>F;632-65/Domain: transmembrane #status predicted <TRR>F;831-65/Domain: transmembrane #status predicted <TRR>F;841-65/Domain: transmembrane #status predicted <TRR
F;841-65/Domain: transmembrane #status predicted <TRR
F;841-65/Domain: transmembrane #status predicted <TRR
F;84
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ï
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 F:131,200,422,498/Binding site: carbohydrate (Asn) (covalent) #status predicted F:593,713/Binding site: phosphate (Ser) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               glutamate receptor delta-1 chain precursor - mouse
C:Species: Mus musculus (house mouse)
C:Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 20-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Species: Methanococcus jannaschii
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 21-Jul-2000
C;Accession: C64396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Gaps
                                                                                                                                                                              A,Molecule type: mRNA
A,Residues: 83-1009 <SPR>
A,Cross-references: BMBL:Z17238; NID:956285; PIDN:CAA78936.1; PID:956286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2; Length 1009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      49.0%; Score 51.5; DB 2; Length 1009;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               precorrin-2 methyltransferase homolog - Methanococcus jannaschii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Indels
R,Sprengel, R.
submitted to the EMBL Data Library, October 1992
A,Reference number: 831222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47.4%; Pred. No. 13; tive 6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     698 PLEQDSTFAELWRTISKNG 716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           49.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  698 PLEQDSTFAELWRTISKNG 716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9; Conservative
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Best Local Similarity
2, Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                           A; Accession: S31222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouerv Match
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Gaps

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GenCore version 5.1.3

Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 12, 2003, 10:32:56; Search time 11 Seconds
71.641 Million cell updates/sec
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Title: US-09-403-440A-2 Perfect score: 105 Sequence: 1 PPLSQETFSDLWKLLPENG 19

Sequence: 1 PPLSQETFSDLWKL: Scoring table: BLOSUM62

rable: BLOSUM62 Gapop 10.0 , Gapext 0.5 Total number of hits satisfying chosen parameters: 112892

112892 segs, 41476328 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : SwissProt\_40:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

	Description	O9tub2 sus scrofa	Q9wur6 cavia porce				P56423 macaca fasc	macaca	Q9ttal tupaia glis	ovis a	Q29628 bos taurus		canis			Q00366 mesocricetu	P07193 xenopus lae			P25035 oncorhynchu	Q9w679 tetraodon m			Q58181 methanococc		bacillu	pepper	pepper					O66651 aquifex aeo	
SOUTHWANTES	QI .	P53 PIG	P53_CAVPO	P53_RABIT	P53_CERAE	P53 HUMAN		P53_MACMU	P53_TUPGB	P53_SHEEP	P53_BOVIN	P53_MARMO	P53_CANFA	P53_FELCA	P53_CRIGR	P53_MESAU	P53_XENLA	P53_MOUSE	P53_RAT	P53_ONCMY	P53_TETMU	P53_XIPHE	P53_XIPMA	Y771_METJA	IPI_BACSU	SLPO_BACBR	COAT PMMV2	COAT_PMMVI	COAT PMMVS	COAT TOMKO		P53_SPEBE	SYI AQUAE	YA24_SYNY3
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	Match Length	386	391	391	393	393	393	393	393	382	386	391	381	386	393	396	363	390	391	396	367	342	342	230	119	1004	156	156	156	156	158	314	926	307
* Query	Match	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	89.5	89.5	89.5	85.7	85.7	81.9	81.9	77.1	70.5	68.6	57.1	51.4	51.0	-	47.6	45.7	n.	4	4	44.8		44.8	4.	44.8	43.8
	Score	66	66	66	66	66	66	66	66	94	94	94	90	90	98	98	91	74	72	9	Ň	٠	53.5		48	48	47	47	47	47	47	47	47	46
Result	No.	-	7	m	4	S	9	7	<b>6</b> 0	6	10	11	12	13	14	15	16	17	18	19	20	21			24		26	27	28	29	30	31	32	33

P52633 mus musculu Q13188 homo sapien	P08343 bovine papi P58879 methanopyru	P05861 simian immu	P11262 simian immu	Q60320 methanococc	P10360 gallus gall	P46207 aquifex pyr	Q00976 kluyveromyc	P39847 bacillus su
STA6_MOUSE STK3_HUMAN	VL3 BPV4 HIS7 METKA	NEF_SIVMK	NEF SIVML	YOOF METUA	P53 CHICK	SYI AQUPY	CC68 KLULA	PPS3_BACSU
		٦.	٠,	Н	Н	Н	ч	-
837	113	207	263	276	367	529	1033	2555
42.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9
45 44.5	4. 4. 4. 4.	4 4	44	44	44	44	44	44
34 35	36 37	38	40	41	42	43	44	45

## ALIGNMENTS

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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELUTAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             D'erchia A.M., Pesole G., Tullo A., Saccone C., Sbisa E.; "Guinea pig p53 mRNA: identification of new elements in coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cavia porcellus (Guinea pig).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Rodentia, Hystricognathi, Caviidae, Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nučlear protein; Phosphorýlation; Apoptosis.
DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC).
DNA BIND 100 290 BY SIMILARITY.
                                                                                                                                                           ..
0
                                                                                                                 94.3%; Score 99; DB 1; Length 386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      untranslated regions and their functional and evolutionary
                                                                                                                                                        0; Indels
318 349 OLIGOMERIZATION.
361 380 BASIC (REPRESSION OF DNA-B 304 316 NUCLEAR LOCALIZATION SIGNA PHOSPHORYLATION (BY SIMILA 386 AA; 42862 MW, A4C3D88EBDF55162 CRC64;
                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-UTN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                    7.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                                    391 AA.
                                                                                                                                100.0%; Preq. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IN MANY TYPES OF CANCER.
-!- SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=99265972; PubMed=10331945;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AJ009673; CAB43196.1; -.
                                                                                                                                                                                              1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                 12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HSSP; P04637; IYCS.
InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genomics 58:50-64(1999).
                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                  Local Similarity
nes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        323
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       implications."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                  P53 CAVPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
                                                                          SEQUENCE
                                                                                                                   Query Match
                                                         MOD RES
                                       OMAIN
                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                 P53_CAVPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The CDMA cloning automicroperson in many tumor types; induces the 185:169-173(1997).

Conclusionated at the control of the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases.

Controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases.

Controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases.

Controlling a set of genes required for this process. One of BAX and FAS antigen expression, or by repression of BCl-2 expression (By similarity).

Controlling binds DNA as a homotetramer (By similarity).

Controlling BIN FORTION: Nuclear.

Controlling BIN FREQUENTLY MUTATED OR INACTIVATED OR INACTIVATED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=97208869; PubMed=9055811; le Goas F., May P., Ronco P., Caron de Fromentel C.; "CDNA cloning and immunological characterization of rabbit p53.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                      orgo.
Orgotolagus cuniculus (Rabbit).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Lagomorpha, Leporidae, Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                massr; Processis 123.
Pfam, PR00870, P53, 1.
PRINTS; PR00386; P53SUPPRESSR.
Prodom, PD002681, P53, 1.
PROSITE, PS00348; P53, 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nuclear protein; Phosphorylation; Apoptosis.

1 43 TRANSCRIPTION ACTIVATION (ACIDIC).
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0
                                                                                                  DB 1; Length 391;
                                                        321D40702383573E CRC64;
                                                                                                                                                                                                                                                                                                                                                           01-NOV-1997 (Rel. 35, Created)
1-NOV-1997 (Rel. 35, Last sequence update)
15-UNV-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                   7.3e-08;
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OLIGOMERIZATION.
                                                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IN MANY TYPES OF CANCER. SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                       Pred. No.
                                                                                                  94.3%; Score 99;
                                                            43287 MW;
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                                                                                                                       100.08;
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                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                          STANDARD;
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                                                                                                                 Local Similarity
hes 18; Conserv
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                                                          391 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9986;
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366
308
390
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SEQUENCE
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393 AA; 43696 MW; 9ED285C9A7855D6E CRC64;
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    SO SEQUENCE
                                                                                                                                                                                                        P53 HUMAN
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                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its most by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC).

DOMAIN 22 356 OLIGOMERIZATION.

DOMAIN 38 387 BASIC (REPRESSION OF DNA-BINDING).

NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).

MOD_RES 392 PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression.
SUBDILI: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISBASE: BS IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. PS3 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                                    Cercopithecus aethiops (Green monkey) (Grivet).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae,
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                                   94.3%; Score 99; DB 1; Length 391; 100.0%; Pred. No. 7.3e-08;
                                                                 0; Indels
      391 AA; 43435 MW; 86BD5B8D0B726525 CRC64;
                                                                                                                                                                                                                                                                                     Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                       01-07N-1990 (Rel. 13, Created)
01-07NV-1990 (Rel. 13, Last sequence update)
15-07UN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                           393 AA.
                                               100.0%; Prec. ...
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MEDLINE=90045967; PubMed=2530498;
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                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                       12 PPLSQETFSDLWKLLPEN 29
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InterPro; IPR002117; P53.
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PROSITE; PS00348; P53; 1.
                                                    Local Similarity 100.
Les 18; Conservative
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         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Harlow E., Williamson N.M., Ralston R., Helfman D.M., Adams T.E.; "Molecular cloning and in vitro expression of a cDNA clone for human
                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=85230577; PubMed=4006916;
Zakut-Houri R., Bienz-Tadmor B., Givol D., Oren M.;
"Human p53 cellular tumor antigen: cDNA sequence and expression in
COS cells.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cellular tumor antigen p53 (Tumor suppressor p53) (Phosphoprotein p53) (Antigen NY-CO-13).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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MEDLINE=87089826; PubMed=3025664;
Harris N., Brill E., Shohat O., Prokocimer M., Wolf D., Arai N.,
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Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
"ps3 is frequently mutated in Burkitt's lymphoma cell lines.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Molecular basis for heterogeneity of the human p53 protein."; Mol. Cell. Biol. 6:4650-4656(1986).
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Matlashewski G., Lamb P., Pim D., Peacock J., Crawford L.,
94.3%; Score 99; DB 1; Length 393; 100.0%; Pred. No. 7.3e-08; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                 P04637; Q16848; Q9UBI2;
13-AUG-1987 (Rel. 05, Created)
01-MAR.1989 (Rel. 10, Last sequence update)
15-UUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                         393 AA
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MEDLINE=87064416; PubMed=2946935;
Lamb P., Crawford L.;
"Characterization of the human p53 gene.";
MOl. Cell. Biol. 6:1379-1385(1986).
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Mol. Cell. Biol. 5:1601-1610(1985).
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MEDLINE=85267676; PubMed=3894933;
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MEDLINE=89108008; PubMed=2905688;
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                                                                                                                            1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                 12 PPLSQETFSDLWKLLPEN 29
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                                                                       18; Conservative
                                                                                                                                                                                                                                                                                                                                         STANDARD;
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         Query Match
Best Local Similarity
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1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                        MEDLINE=96197761; PubMed=8632903; Flaman J.-M., Waridel F., Estreicher A., Vannier A., Limacher J.-M., Gilbert D., Iggo R., Frebourg T.; "The human tumour suppressor gene p53 is alternatively spliced in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=95292092; PubMed=7773777;
Leé W., Harvey T.S., Yin Y., Yau P., Litchfield D., Arrowsmith C.H.;
"Solution structure of the tetrameric minimum transforming domain of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=94294808; PubMed=8023159;
Clore G.M., Omichinski J.G., Sakaguchi K., Zambrano N., Sakamoto H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shaw P., Freeman J., Bovey R., Iggo R., Regulation of specific DNA binding by p53: evidence for a role for O-G1ycosylation and charged residues at the carboxy-terminus."; Oncogene 12:921-930(1996).
                                                                                                                                                Anderson C.W., Kieleczawa J., Allalunis-Turner J.; "Human p53 from the malignant glioma-derived cell lines M059J and M059K have a cancer-associated mutation in exon 8."; Submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Appella B., Gronenborn A.M.; "High-resolution gruncture of the oligomerization domain of p53 by
                                                                                                                                                                                                                                                                                                                                       Addison C., Jenkins J.R., Sturzbecher H.-W.;
"The p53 nuclear localisation signal is structurally linked to a p34cdc2 kinase motif.";
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Bischoff J.R., Friedman P.N., Marshak D.R., Prives C., Beach D
"Human p53 is phosphorylated by p60-cdc2 and cyclin B-cdc2.";
Proc. Natl. Acad. Sci. U.S.A. 87:4766-4770(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                           Hong T.M., Chen J.J., Peck K., Yang P.C., Wu C.W., "p53 amino acids 339-346 represent the minimal p53 repression
                                     Chumakov P.M., Almazov V.P., Jenkins J.R.;
Submitted (JUN-1991) to the EMBL/GenBank/DDBJ databases.
                                                                                 Rozemuller E.H., Tilanus M.G.J.;
"P53 genomic sequence. Corrections and polymorphism.";
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
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Biol. Chem. 276:1510-1515(2001).
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Nat. Struct. Biol. 1:877-890(1994).
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MEDLINE=94294808; PubMed=8023159;
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                                                                                                                                                                                                                                                                                                                 NUCLEAR LOCALIZATION SIGNAL
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  EMBO J. 10:2879~2887 (1991),
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                                                                                                                                                                                                                                                                                                                                                                                                     MINIMAL REPRESSION DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                               Oncogene 5:423-426(1990).
                                                                                                                                                                                                               ALTERNATIVE SPLICING.
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                       SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                            normal cells.";
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                      MEDILINE-98026899, PubMed-3321402, MCCOy M., Stavridi E.S., Waterman J.L., Wieczorek A.M., Opella S.J., Halazonetis T.D., "Hydrophobic side-chain size is a determinant of the three-dimensional structure of the p53 oligomerization domain.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kussie P.H., Gorina S., Marechal V., Elenbaas B., Moreau J.,
Levine A.J., Pavletich N.P.,
"Structure of the MDM2 oncoprotein bound to the p53 tumor suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=96271983; PubMed=8829653; de Vries T.N., Hartmann A., Blaszyk H., Liao D., Soussi T., Kovach J.S., Sommer S.S.; "Database of mutations in the p53 and APC tumor suppressor genes designed to facilitate molecular epidemiological analyses."; Hum. Mutat. 7:202-213(1996).
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MEDLINE=97081050; PubMed=8875929;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Olschwang S., Laurent-Puig P., Vassal A., Salmon R.-J., Thomas G., "Characterization of a frequent polymorphism in the coding sequence of the Tp53 gene in colonic cancer patients and a control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=95184011; PubMed=7878469; Jeffrey P.D., Gorina S., Pavletich N.P.; Crystal structure of the tetramerization domain of the p53 tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
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                                                                                                                                                                                                                                                                                                                   MEDLINE=94294806; PubMed=8023157; Cho Y., Gorina S., Jeffrey P.D., Pavletich N.P.; Cho Y., Gorina S., Jeffrey P.D., Pavletich N.P.; "Crystal structure of a p53 tumor suppressor-DNA complex: understanding tumorigenic mutations."; Science 265:346-355(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REVIEW ON VARIANTS.
MEDLINE=91289156; PubMed=1905840;
Hoolstein M., Sidransky D., Vogelstein B., Harris C.C.;
"p53 mutations in human cancers.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CRYSTALLOGRAPHY (1.7 ANGSTROMS) OF 325-356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'Match 94.3%; Score 99; DB 1; Le Local Similarity 100.0%; Pred. No. 7.3e-08; les 18; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                        X-RAY CRYSTALLOGRAPHY (2.2 ANGSTROMS) OF 94-289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=94090335; PubMed=8266092;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=91153807; PubMed=1999338;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       suppressor at 1.7 angstroms.";
Science 267:1498-1502(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genet. 86:369-370(1991).
STRUCTURE BY NMR OF 326-354.
                                                                                                                                                                                                                    EMBO J. 16:6230-6236(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Science 274:1001-1005(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Science 262:1980-1981(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Science 274:948-953(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transactivation domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ience 253:49-53(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REVIEW ON VARIANTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          VARIANT ARG-72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harris C.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        population.",
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                                                                                                                                                                                                                                                                                                                                Khan M.A. Hansen C., Welsh J.A., Bennett W.P.;
SEQUENCE FROM N.A.

Khan M.A., Hansen C., Welsh J.A., Bennett W.P.;
Submitted (FEB-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                          Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BY SIMILARITY.
OLIGOMERIZATION.
BASIC (REPRESSION OF DNA-BINDING).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nuclear protein, Phosphorylation, Apoptosis.

DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PHOSPHORYLATION (BY SIMILARITY)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.3%; Score 99; DB 1; Length 393; 100.0%; Pred. No. 7.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUCLEAR LOCALIZATION SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EDEEBA1BD3AEAFB8 CRC64;
                                                                                                                                        15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prec. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43678 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; U48957; AAB91535.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MANY TYPES OF CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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InterPro; IPR002117; P53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                   Cercopithecinae; Macaca.
                                                                                                         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         325
368
311
392
393 AA;
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                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9541;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HSSP; P04637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
                                                                                                                                                                                                             TP53 OR P53.
                                                                                                       P53 MACFA
P56423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
MOD RES
SEQUENCE
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                                                                                         P53_MACFA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Khan M.A., Hansen C., Welsh J.A., Bennett W.P.;
Khan M.A., Hansen C., welsh J.A., Bennett W.P.;
Khan M.A., Hansen C. to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces

-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces

-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces

-!- FUNCTION: Acts or apoptosis depending on the physiological

-!- FUNCTION: Acts or apoptosis depending on the physiological

-!- FUNCTION: Acts or apoptosis dependent cell division

by controlling a set of genes required for this process. One of

the activated genes is an inhibitor of cyclin-dependent kinases.

Apoptosis induction seems to be mediated either by stimulation of

BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE-Blood;
MEDLINE-9411042; PubMed-8125305;
KAY H.D., Mountjoy C.P., Wu G., Cornish K.G., Smith L.J.;
"Sequence of a cDNA encoding the p53 protein in rhesus monkey (Macaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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SUBGELLULAR LOCATION: Nuclear.
DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                           Macaca milatta (Rhesus macaque).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BASIC (REPRESSION OF DNA-BINDING). NUCLEAR LOCALIZATION SIGNAL. PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PERMY PR00870; P53; I. PRINTS; PR0386; P53STPRESSR.
ProDom, P0002681; P53; I. PROSITE; P500348; P53; I. Anti-oncogene; DNA-binding; Transcription regulation; Activator Nuclear protein; Phosphorylation; Apoptosis.

| Anti-oncogene; Phosphorylation; Apoptosis. | Anti-oncogene; Phosphorylation; Apoptosis. | Anti-oncogene; Phosphorylation; Apoptosis. | Anti-oncogene; Phosphorylation (ACIDIC).
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Pred. No. 7.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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                                                                                                                                           Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BY SIMILARITY.
OLIGOMERIZATION.
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100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                annotation update)
393 AA
                                                                                                 Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          43655 MW;
                                                                  15-JUL-1998 (Rel. 36, Created)
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EMBL; U48956; AAB91534.1; -.
HSSP; P04637; lOLG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IN MANY TYPES OF CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Watches 18; Conservative
                                                                                                    (Rel. 36, Last
(Rel. 41, Last
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR002117; P53.
   STANDARD;
                                                                                                                                                                                                                                                                                                                                       Cercopithecinae, Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mulatta).";
Gene 138:223-226(1994).
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                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expression
                                                                                                    15-JUL-1998
15-JUN-2002
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DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression.
--- SUBUNT: Binds DNA as a homotetramer (By similarity).
--- SUBCELLULAR LOCATION: Nuclear.
--- SUBCELLULAR LOCATION: Nuclear.
--- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                  Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Scandentia, Tupaiidae, Tupaia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nuclear protein; Phosphorylation; Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 393;
                                                  P53_TUPGB STANDARD; PRT; 393 AA. 09TTA1; CCT-2001 (Rel. 40, Created) 16-OCT-2001 (Rel. 40, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) 16-JUN-2002 (Rel. 41, Last annotation update) 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OLIGOMERIZATION.
                                                                                                                                                                                                                                                                Tupaia glis belangeri (Common tree shrew).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=Chinensis; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AF175893; AAF22640.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00386; P53SUPPRESSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43552 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HSSP; P04637; IYCS.
InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               393 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9396;
                                                                                                                                                                                                                                          OR P53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOMAIN
MOD RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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RESULT 8
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BASIC (REPRESSION OF DNA-BINDING).
NUCLERA LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
FD9F36603945A1FA CRC64,
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0;
                     Gaps
                     ;
0
                     0; Indels
Score 99; DB 1; Losted. No. 7.3e-08;
    100.0%; Pred. w.
        Best Local Similarity 100.
Matches 18; Conservative
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ò QQ RESULT 10 P53 BOVIN

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1 PPLSQETFSDLWKLLPEN 18
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RESULT 9

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between the Swiss Institute of Bioinformatics and the EMBL outstation.

the European Bioinformatics Institute. There are no restrictions on its
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or send an email to license@lsb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration
                                                                                                                                                                                                                                                                                                                                   DIA Seq. 5:255-259(1995).

-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of bax and FAS antigen expression, or by repression of BCl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                              Dequiedt F., Kettmann R., Burny A., Willems L.; "Nucleotide sequence of the ovine P53 tumor-suppressor cDNA and its genomic organization.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SUBUNIT: Binds DAT ALC. -!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                               Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                          Ovis aries (Sheep).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ProDom: PRO02681; P53; 1.
PROSITE; PRO0348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
Nuclear protein; Phosphorylation; Apoptosis.
TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression. SUBUNIT: Binds DNA as a homotetramer (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89.5%; Score 94; DB 1; Lengtn 302. 94.4%; Pred. No. 4.2e-07; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4B024076C79C3B2D CRC64;
                                                  01-OCT-1996 (Rel. 34, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OLIGOMERIZATION.
   382 AA
                                                                                                                                                                                                                                                                     MEDLINE=95352828; PubMed=7626788;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PR00386; P53SUPPRESSR.
                                   01-OCT-1996 (Rel. 34, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42809 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; X81705; CAA57349.1; -. HSSP; P04637; 1C26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                              Caprinae; Ovis.
 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 17; Conservative
                                                                                                                                                             Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         382 AA;
                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                               NCBI_TaxID=9940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    314
357
300
381
                                                                                                                                                                                                                                                      rissue=Blood
P53_SHEEP
P51664;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOMAIN
DNA BIND
DOMAIN
                                                                                                                                                                                Bovidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPECIES-B. indicue; STRAIN-Boran; TISSUE-Blood;
Bishop R.R.P., Gobright B.E.I.;
Bishop R.R.P., Gobright B.E.I.;
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.

-I-FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and call type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Komori H., Ishiguro N., Horiuchi M., Shinagawa M., Aida Y.;
"Predominant p33 mutations in enzootic bovine leukemic cell lines.";
Vet. Immunol. Immunopathol. 52:53-63(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISEASE: P53 IS FOUND IN INVERASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                         Bos indicus (Zebu).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-oncogene; DNA-binding; Transcription regulation; Activator; Nuclear protein; Phosphorylation; Apoptosis.
                                                                                                                                                                                                                                                                                           SPECIES-Bovine; TISSUE-Liver;
MEDLINE-9935-28429; PubMed=7626789;
Dequiedt F., Kettmann R., Burny A., Willems L.;
"Nucleotide sequence of the bovine P53 tumor-suppressor CDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TRANSCRIPTION ACTIVATION (ACIDIC)
                                           01-NOV-1997 (Rel. 35, Created)
1-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 13-386 FROM N.A.
SPECIES=Bovine; STRAIN=Holstein; TISSUE=Thymus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BY SIMILARIT
                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=96401400; PubMed=8807776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL, X81704; CAA57348.1; -. EMBL; D49825; BAA08629.1; -. EMBL; U74486; AAB51214.1; -. HSSP; P04637; ITUP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                     DNA Seq. 5:261-264(1995).
                                                                                                                                         Bos taurus (Bovine), and
         STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                285
349
316
385
                                                                                                                                                                                                               Bovidae, Bovinae, Bos.
NCBI_TaxID=9913, 9915;
                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA BIND
DOMAIN
P53 BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DOMAIN
MOD_RES
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NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BGl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-97376996; PubMed=9233767; Feitelson M.A., Zhang S.M.; Feitelson M.A., Ranganathan P.N., Clayton M.M., Zhang S.M.; "Partial characterization of the woodchuck tumor suppressor, p53, and its interaction with woodchuck hepatitis virus X antigen in hepatocarcinogenesis."; Oncogene 15:327-336(1997).
                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISEAGE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                        Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Sciuridae; Sciurinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nuclear protein; Phosphorylation; Apoptosis.

DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC).
DNA_BIND 100 290 BY SIMILARITY.

DOMAIN 323 354 OLIGOMERIZATION.
                                                                                             ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBUNIT: Binds DNA as a homotetramer (By similarity). SUBCELLULAR LOCATION: Nuclear.
                                                        Length 386;
                                                                                             1; Indels
380 380 R -> T (IN REF. 2).
386 AA; 43255 MW; 222473F28C548F31 CRC64;
                                                                                                                                                                                                                                                                                                                                               15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                            Pred. No. 4.3e-07;
                                                        89.5%; Score 94; DB 1; 94.4%; Pred. No. 4.3e-07
                                                                                                                                                                                                                                                                       PRT; 391 AA
                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                 Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AJ001022; CAA04478.1; -. HSSP; P04637; 1TUP.
InterPro; IPR002117; P53.
Pfam; PR00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; P0002681; P53; 1.
                                                                                                                                                                                                                                                                                                             16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IN MANY TYPES OF CANCER.
                                                                                                                                     1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                   12 PPLSQETFSDLWNLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                            Marmota monax (Woodchuck)
                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                         STANDARD;
                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression.
                                                                                                                                                                                                                                                                           P53 MARMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DOMAIN
MOD RES
                        SEQUENCE
      CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Marmota
                                                                                                                                                                                                                                                                                             036006;
                                                                                                     Matches
                                                                                                                                                                                                                                      RESULT 11
P53_MARMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Setoguchi A., Sakai T., Okuda M., Minehata K., Yazawa M., Ishizaka T.,
Watari T., Hasagawa A., Tsujimoto H.,
"Aberrations of p53 tumor suppressor gene in various spontaneous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-Leukocyte;
MEDLINE-SB178696; PubMed=9519881;
Veldhoen N., Milner J.;
"Isolation of canine p53 cDNA and detailed characterization of the full length canine p53 protein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Canis familiaris (Dog).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Carnivora, Fissipedia, Canidae, Canis.
                                                                                                        ;
                                            89.5%; Score 94; DB 1; Length 391; 94.4%; Pred. No. 4.3e-07; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases.
391 AA; 43468 MW; EIDESDB84BA40182 CRC64;
                                                                                                                                                                                                                                                                                                                   P53 CANPA STANDARD; PRT; 381 AA. 0129537; Q9TV78; Clasted) 15-DEC-1998 (Rel. 35, Created) 15-DEC-1998 (Rel. 37, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; AF060514; AAC16909.1; -. EMBL; AR020761; BAA78379.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 25-300 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IN MANY TYPES OF CANCER.
                                                                                                                                             1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                            12 PPLSQETFSDLWNLLPEN 29
                                                                                                   Conservative
                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              dog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rissum=spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN=Beagle;
                                                                                                   17;
SEOUENCE
                                                                                              Matches
                                                                                                                                                                                                                                                                   RESULT 12
                                                                                                                                                                                                                                                                                              P53_CANFA
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BY SIMILARITY.

OLIGOMERIZATION.

BASIC (REPRESSION OF DNA-BINDING).

NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).

PHOSPHORYLATION (BY SIMILARITY).

MES. > MCGP (IN REF. 2).

L. > P (IN REF. 2).

MW; 761A718FDC93DA59 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     J. vet. Med. Sci. 55:801-805(1993).

-! FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bc1-2
                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUB=Lymph node,

MEDLINE=5433360; PubMed=8056458;

Okuda M., Umeda A., Sakai T., Ohashi T., Momoi Y., Youn H.Y.,

Watari T., Goitsuka R., Tsujimoto H., Hasegawa A.;

"Cloning of feline p53 tumor-suppressor gene and its aberration in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=94114699; PubMed=8286534; Okuda M., Umeda A., Matsumoto Y., Momoi Y., Watari T., Goitsuka R., O'Srien S.J., Tsujimoto H., Hasegawa A.; "Molecular cloning and chromosomal mapping of feline p53 tumor suppressor gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SUBDATT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: PS3 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
--- DISEASE: PS3 IS FREQUENTLY MUTATED OR INACTIVATED IN MANY TYPES OF CANCER.
-!- SIMILARITY: BELONGS TO THE PS3 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Felis silvestris catus (Cat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
                                                                                                                   Anti-oncogene, DNA-binding, Transcription regulation, Activator,
                                                                                                                                                                                                                                                                                                                                                              ·,
                                                                                                                               Nuclear protein; Phosphorylation; Apoptosis.
DOMAIN 44 TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                   85.7%; Score 90; DB 1; Length 381;
88.9%; Pred. No. 1.7e-06;
.ive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-vUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
TP53 OR TRP53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                386 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Last sequence update)
              HSSP, P04637; 10LG.
InterPro; IPR002117; P53.
Pfam; P700870; P53; 1.
PRINTS; PR00386; P533UPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  int. J. Cancer 58:602-607(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequ
15-JUN-2002 (Rel. 41, Last anno
                                                                                                                                                                                                                                                                                          42486 MW;
EMBL; S77819; AAB42022.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 34-354 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                         1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                        12 PPLSQETFSELWNLLPEN 29
                                                                                                                                                                                                                                                                                                                                                         16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STANDARD;
                                                                                                                                                                                     344
                                                                                                                                                                                                                                    380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hematopoietic tumors
                                                                                                                                                                                                                                                                                      381 AA;
                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                   313
356
299
380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9685;
                                                                                                                                                                     8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P53 FELCA
                                                                                                                                                                   DNA_BIND
                                                                                                                                                                                                                                                 CONFLICT
                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                    RES
                                                                                                                                                                                                                                                                      CONFLIC
                                                                                                                                                                                   DOMAIN
                                                                                                                                                                                                    DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P41685;
                                                                                                                                                                                                                      DOMAIN
                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ·;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

TISSUE-Embryonic fibroblast;
Shimizu T., Nikaido O., Suzuki F.;
Submitred (JUN-1996) to the EMBL/GenBank/DDBJ databases.

Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.

In FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                 BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
                                                                                                                                                                                                    InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00346; P53SUPPRESSR.
ProDom; P0002681; P53; 1.
PROSITE; P800348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
                                                                                                                                                                                                                                                                                                              Nuclear protein, Phosphorylation, Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lee H., Larner J.M., Hamlin J.L.; "Cloning and characterization of Chinese hamster p53 cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.7%; Score 90; DB 1; Length 386; 88.9%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         i, Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chaung W., Mi L.J., Boorstein R.J.;
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                      K -> R (IN REF. 2).
D08B43BA1BC8EB78 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               009185; Q64397; P97288; P97788; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) Cellular tumor antigen p53 (Tumor suppressor p53). TP53 OR P53.
                                                                                                                                                                                                                                                                                                                                                BY SIMILARITY.
OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      393 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cricetulus griseus (Chinese hamster).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=97183659; PubMed=9031625;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        42692 MW;
                                                                                                                                                   EMBL; D26608; BAA05653.1; -. EMBL; D16460; BAA03927.1; -. HSSP; P04637; IOLG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 PPLSQETFSELWNLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gene 184:177-183(1997).
                                                                                                                                                                                                                                                                                                                                                                                   361
304
385
385
285
386 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10029;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rissum=Liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cricetulus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P53 CRIGR
                                                                                                                                                                                                                                                                                                                              DOMAIN
DNA BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                 DOMAIN
DOMAIN
MOD RES
CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 14
P53_CRIGR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN=Syrian; TISSUB=Kidney;
MEDLINE=92210007; PubMed=155573;
Legros Y., McIntyre P., Soussi T.;
"The cDNA cloning and immunological characterization of hamster p53.";
Gene 112:247-250(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BY SIMILARITY.
OLIGOMERIZATION
BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
L -> Q (IN CELL LINE V79-4).
Y -> F (IN REF. 2).
Apoptosis induction seems to be mediated either by stimulation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                          SUBGINIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISEASE: F53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. F53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nuclear protein; Phosphorylation; Apoptosis.
DOMAIN 45 TRANSCRIPTION ACTIVATION (ACIDIC)
              BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                81.9%; Score 86; DB 1; Length 393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -> F (IN REF. 2).
2A7830E788311689 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. 7.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PR3 MESAU STANDARD; PRT; 396 AA. 000366; P97276; 01-DEC-1992 (Rel. 24, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                          SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mesocricetus auratus (Golden hamster).
                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P538UPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 43378 MW;
                                                                                                                                                                                                                                                                            EMBL; Y08900; CAA70108.1; -.
EMBL; Y08901; CAA70109.1; -.
EMBL; U50395; AAC53040.1; -.
EMBL; D86070; BAA13004.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13 PLSQETFSDLWKLLPPN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                133
135
103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       102
325
368
368
391
133
103
393 AA;
                                                                                                                                                                                                                                                                                                                                               HSSP; P04637; 1TUP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            [2]
SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=10036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA BIND
DOMAIN
DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOD RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VARĪANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 15
P53_MESAU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
BY SIMILARITY.
OLIGOMERIZATION.
DASJIC (FEPRESSION OF DNA-BINDING).
NUCLERA LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                expression.
-!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSPORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
-!- SIMILARITY: BELONGS TO THE P53 PAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ProDon; PROUSE; PS3; 1.
PROSITE; PS00268; PS3; 1.
PROSITE; PS00368; PS3; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
Nuclear protein; Phosphorylation; Apoptosis.
TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81.9%; Score 86; DB 1; Length 396; 94.1%; Pred. No. 7.5e-06; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G -> S (IN REF. 2).
906EF02568099BE3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                            EMBL; M75144; AAA37085.1; -. BMBL; U07182; AAB41344.1; -. PIR; JH0633; JH0633. HSSP; P04637; 1TUP. InterPer; IRFO0870; P53; PFam; PF00870; P53; 1. PRINTS; PR00386; P53SUPPRESSR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         43631 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               314
395
395
188
196 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
DNA_BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOD RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DOMAIN
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Search completed: February 12, 2003, 10:35:02 Job time : 12 secs

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February 12, 2003, 10:33:27; Search time 29 Seconds (without alignments) 134.996 Million cell updates/sec
GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                     671580 segs, 206047115 residues
                                                                                 OM protein - protein search, using sw model
                                                                                                                                                                                                             US-09-403-440A-2
105
1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                          Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                      BLOSUM62
                                                                                                                                                                                                                                    Perfect score:
                                                                                                                                                                                                                                                                                                      Scoring table:
                                                                                                                                                                                                                                                             Sequence:
                                                                                                                                                                                                                                                                                                                                                                     Searched:
                                                                                                                            Run on:
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671580

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Minimum DB seq length: 0 Maximum DB seq length: 200000000

sp archea:\*
sp archea:\*
sp fungi:\*
sp human:\*
sp invertebrate:\*
sp mammal:\* sp\_unclassified:\* sp\_rodent:\*
sp\_virus:\*
sp\_vertebrate:\* sp\_rvirus:\* sp\_bacteriap:\* sp\_organelle:\* sp\_phage:\* sp plant:\* SPTREMBL 111: 123: 14: 15: 16: Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

sp\_archeap:\*

			Description	Q9nzd0 homo sapien	Q8spz3 delphinapte	Q9uq61 homo sapien	homo	Q16811 homo sapien	homod	homo	homo	homo	Q16808 homo sapien	Q16809 homo sapien	Q16810 homo sapien	homo	Q16535 homo sapien	Q9n252 sus scrofa	Q920y0 meriones un
SUMMARIES			ID	O9NZD0	Q8SPZ3	Q9UQ61	Q9HAQ8	Q16811	Q16807	Q15086	Q15087	Q15088	Q16808	Q16809	016810	Q9BTM4	Q16535	Q9N252	1 Q920Y0
		í	2	4	9	4	4	4	4	4	4	4	4	4	4	4	4	9	11
	de ,		Length	32	387	393	393	393	393	393	393	393	393	393	393	393	393	387	390
	æ (	Query	March	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	89.5	83.8
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77 73.3 25 4 Q9NP68 74 70.5 397 11 Q9ER40 74 70.5 397 11 Q9ER40 72 68.6 271 11 Q9EEP9 69 67.1 265 13 Q9W681 60 57.1 265 13 Q9W681 60 57.1 265 13 Q9W682 52 49.5 443 5 Q9WG7 51.5 49.0 1009 11 Q62640 51.5 49.0 1009 11 Q6241 50 47.6 367 16 Q9ER57 49 46.7 16 Q9TW4 49 46.7 16 Q9TW4 41.8 157 12 Q91U97 47 44.8 157 12 Q91U97 47 44.8 158 12 Q91U97
7 7 7 7 7 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
78 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4

### ALIGNMENTS

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CTISSUB-GILOMAS AND ASSOCIATED NORMAL TISSUES;
TISSUB-GILOMAS AND ASSOCIATED NORMAL TISSUES;
TISSUB-GILOMAS AND ASSOCIATED NORMAL TISSUES;
Thompson-Hehir J., Davises M.P.A., Green J.A., Halliwell N.,
Joyce K.A., Salisbury J., Sibson D.R., Vergote I., Walker C.;
Thuration Detection Utilizing a Novel PCR Approach for Amplification
of the p53 Gene from Microdissected Tissue: Application to Archival
Thuror Samples.";
Submitted (DEC-1999) to the EMBL/GenBank/DDBJ databases.
REMBL, AF210308; AAR63442.1; -
REMBL, AF210308; AAR63442.1; -
REMBL, AF210308; AAR63442.1; JOINED.
REMBL, AF210308; AAR63442.1; JOINED.
REMPL, AF210308; AAR63442.1; JOINED.
REPERPROPORTOR P53; 1.
ProDom, PD002681; P53; 1.
PRODOM; PD002681; P53; 1.
REMDL TERM REMBL AF34040.1; DNN TERM REMDL REM
                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                 01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                                                              32 AA.
                                                                                                                                                                                                                                                                    P53 tumor suppressor protein (Fragment). TP53.
                                                                          PRT;
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Matches 18; Conservative
                                                                              PRELIMINARY;
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RESULT 1
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                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                               Xu N., Shiraki T., Yamada T., Nakajima M., Gauthier J.M.,
Pfeiffer C.J., Sato S.;
"Nucleotide sequence of the p53 cDNA of beluga whale (Delphinapterus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anderson C.W., Kieleczawa J., Allalunis-Turner M.J.; "Human p53 from malignant glioma-derived M059J and M059K cells has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Isolation of two cell lines from a human malignant glioma specimen differing in sensitivity to radiation and chemotherapeutic drugs."; Radiat. Res. 134:349-354(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).
DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF
TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED IN
                                                                                                                                         Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti;
Monodontidae; Delphinapterus.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                                                                                                                                                                                                                   Score 99; DB 6; Length 387;
Pred. No. 9e-08;
Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                        EMBL; AF475081; AAL83290.1; -.
SEQUENCE 387 AA; 43033 MW; E4C8BEDDF34A540E CRC64;
                                                      01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                             PRT; 387 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT; 393 AA.
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94.3%; Score 99; DB
Best Local Similarity 100.0%; Pred. No. 9e-
Matches 18; Conservative 0; Mismatches
                                                                                                                                Delphinapterus leucas (Beluga whale).
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                                                                                                                                                                                                                                                                                                                                                                                                                             12 PPLSQETFSDLWKLLPEN 29
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                             PRELIMINARY;
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                                            Q8SPZ3;
                             O8SPZ3
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RESULT 2
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                  Q8SPZ3
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"Hyaluronidase Induction of a WW Domain-Containing Oxidoreductase that
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1 Biol. Chem. 276:3361-3370(2001).

1 EIO. Chem. 276:3361-3370(2001).

1 FUNCTION: ACTS AS ATUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APPOPTOSIS DEPENDING ON THE PHYSIOLOGICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANS-ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE CELL DIVISION BY CONTROLLING A SET OF GENER REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITIOR OF CELLIN-DEPENDENT KINAGES. APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF
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Probom; P0020881; P53; 1.
PROSITE; PS00348; P53; 1.
Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
MEDLINE=21264809; PubMed=11058590;
Chang N.-S., Pratt N., Heath J., Schultz L., Sleve D., Carey G.B.,
Zevotek N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 99; DB 4; Length 393;
Pred. No. 9.2e-08;
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U-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Cellular tumor antigen p53.
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100.0%; Pred. No. ...
0; Mismatches
MANY TYPES OF CANCER (BY SIMILARITY).

-! SIMILARITY: BELONGS TO THE P53 FAMILY.
EMBL; AF135121; AAD28535.1; --
EMBL; AR135120; AAD28535.1; JOINED.
HSSP; P04637; 10LG.
INTERPLO; IPRO02117; P53.
INTERPLO; IPRO0217; P53.
PEAN; PRO0870; P53; 1.
PRINTS; PR00386; P533UPPRESSR.
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SEQUENCE 393 AA; 43712 MW; AC611E4938C
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Matches 18; Conservative
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Pfam; PF00870; P53; 1.
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A Lamb P., CrawGord L.;

Characterization of the human p53 gene.";

Mol. Cell. Biol. 6:1379-1385(1986).

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES CROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSICLOCAICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANSCOTT TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANSCOTT TOWN SUPPRESSION BY ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION OF BCL-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                     "Isolation and characterization of a human p53 cDNA clone: expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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DISERASE: PS3 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. PS3 IS FREQUENTLY. WITATED OR INACTIVATED IN MANY TYPES OF CANCER (BY SIMILARITY).
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                                                                                                                                                                                                                                                                                Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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94.3%; Score 99; DB 4; Length 393;
Best Local Similarity 100.0%; Pred. No. 9.2e-08;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE-85126934; PubMed-6396087;
Matlashewski G., Lamb P., Pim D., Peacock J., Crawford L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Phosphorylation; Transcription regulation.
SEQUENCE 393 AA; 43698 MW; AD1979573AFALDA5 CRC64;
                                                                                                                                                                     01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                  393 AA
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                                                                                                                                    PRT;
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EMBL; M13112, AAA59987.1; JOINED.
EMBL; M13113; AAA59987.1; JOINED.
EMBL; M13114; AAA59987.1; JOINED.
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M13119; AAA5987.1; JOINED.
M13120; AAA59987.1; JOINED.
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ProDom; PD002681; P53; 1.
                                                                                                                                                                                                                                Cellular tumor antigen p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the human p53 gene.";
EMBO J. 3:3257-3262(1984).
                         12 PPLSQETFSDLWKLLPEN 29
     1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                    PRELIMINARY;
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EMBL;
EMBL;
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EMBL;
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                                                                                                RESULT 5
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SEQUENCE FROM N.A.

WEDLINE=22007731; PubMed=1915267;

REDLINE=22007731; PubMed=23007731;

REDLINE=22007731;

REDLINE
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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100.0%; Pred. No. 9.2e-08;
tive 0; Mismatches 0; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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Last annotation update)
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VARIANT 193 193 R -> H.
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                                                                                                                                                                                                                                                                                   Created)
                                                                                                                                                                                                            PRT;
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                                     12 PPLSQETFSDLWKLLPEN 29
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Homo sapiens (Human).
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Best Local Similarity
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RN FORDENCE FROM N.A.

RA FARTELL P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;

RA FARTELL P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;

RA FARTELL P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;

RI FERD J. 10:2879-2887(1991).

CC G. 10:2879-2887(1991).

CC -!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSIOLOGICAL

CC GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSIOLOGICAL

TUMOR SUPPRESSION. ACTS IN GELL CYCLE REGULATION, IS A TRANS-

CC CONTROLLING A SET OF GENES REGULATE CELL DIVISION BY

CC CONTROLLING A SET OF GENES REQUIRANT ENCLESS. ONE OF THE

ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF

CC BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2

EXPRESSION (BY SIMILARITY).

CC -:- DISABASE F93 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF

TRANSFORMED CELLS. P53 IS FREQUENTLY, WUTATED OR INACTIVATED IN

CC -:- SIMILARITY: BELONGS (BY SIMILARITY).

CC -:- SIMILARITY: BELONGS (BY SIMILARITY).

CC -:- SIMILARITY: BELONGS (BY SIMILARITY).

CC -:- SIMILARITY: BELONGS (BY SIMILARITY).
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                               SEQUENCE FROM N.A.
MEDLINE=9200731; PubMed=1915267;
Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
"p53 is frequently mutated Burkitt's lymphoma cell lines.";
EMBL S601:879-2887(1991).
EMBL, X6001:6; CAA42631.1; -.
                                                                                                                                                                                                                                                                                                                                             94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; Live 0; Mismatches 0; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Cellular tumor antigen p53 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT; 393 AA.
                                                                                                                                                                                                                                                                             Y -> C.
                                                                                                                                                                                                   InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
VARIANT 238 238 Y
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Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; P0002681; P53; 1.
PROSITE; PS00348; P53; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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SEQUENCE
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

MEDLINE-92007731; PubMed=1915267;

Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;

"D53 is frequently mutated Burkitt's lymphoma cell lines.";

EMBD J. 10:2879-2887(1991).

EMBL, X60013; CAA42628.1; -.

HSSP; P04637; 10LG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=92007731; PubMed=1915267; Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.; "p53 is frequently mutated Burkitt's lymphoma cell lines."; BMBO J. 10:2879-2887(1991).
BMBL; X60014; CAA42629.1; -.
HSSP; P04637; 10LG.
                                                                                                                                                                                                                                                                                Query Match 94.3%; Score 99; DB 4; Length 393; Best Local Similarity 100.0%; Pred. No. 9.2e-08; Matches 18; Conservative 0; Mismatches 0; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
P53 transformation suppressor (Fragment).
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last_annotation update)
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                                                                                                                                      InterPro; IRR002117, P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
VARIANT 246 246 T
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Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
VARIANT 237 1.
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SEQUENCE
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Q15087;
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Homo sapiens (Human)
                               Homo sapiens (Human).
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NON TER
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C. 1- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIE DEPENDING ON THE PHYSIOLOGICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, 1S A TRANSCONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVITED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2
EXPRESSION (BY SIMILARITY).

--- SUBSCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).

--- SIBSCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).

--- SIBSCELLULAR TYPES OF CANCER (BY SIMILARITY).

--- SIMILARITY: BELONGS TO THE P53 PAMILY.

REMBL; X60019; CDAA2234.1; ---

--- SIMILARITY: BELONGS TO THE P53 PAMILY.
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                                                  Gaps
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                                                                                                                                                                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
MEDLINE-92007731; PubMed=1915267;
MEDLINE-92007731; PubMed=1915267;
MEDLINE-92007731; PubMed=1915267;
"p53 is frequently mutated Burkitt's lymphoma cell lines.";
EMBO J. 10:2879-2887(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 94.3%; Score 99; DB 4; Length 393; Local Similarity 100.0%; Pred. No. 9.2e-08; es 18; Conservative 0; Mismatches 0; Indels
                      94.3%; Score 99; DB 4; Length 393;
100.0%; Pred. No. 9.2e-08;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      393 AA; 43684 MW; 3DB21E4A398CD0C7 CRC64;
 393 AA; 43627 MW; 3325FAE63F8E1E61 CRC64;
                                                                                                                                                                                      Created)
Last sequence update)
Last annotation update)
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Last sequence update)
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                                                                                                                                                                393 AA.
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                                                                                                                                                                                                                           Cellular tumor antigen p53 (Fragment).
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                                                                                                                                                              PRT;
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Pfam; PF00870; P53; I.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; P0002681; P53; 1.
PROSITE; PS00348; P53; 1.
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                                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01,
                                                                                                                                                                                                     (TrEMBLrel. 01,
                                                                                                                                                                                                                 01-MAR-2002 (TrEMBLrel. 20,
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                                                                           1 PPLSQETFSDLWKLLPEN 18
                                                                                           12 PPLSQETFSDLWKLLPEN 29
                                       Local Similarity Local Similarity Local 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 213
                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                    Homo sapiens (Human)
                                                                                                                                                                                                                                                                                    NCBI_TaxID=9606;
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01-NOV-1996 (
01-NOV-1996 (
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   SEQUENCE
                              Query Match
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Strausberg R.; Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSIOLOGICAL CIRCUMSTANCES OR CELL TYPE, BOT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Activator; Anti-oncogne; Apoptosis; DNA-binding; Nuclear protein;
Phosphorylation; Transcription regulation.
VARIANT 254 254 D -> N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                       Eukaryota, Metazoa; Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          393 AA; 43714 MW; AC69737F2CF9C53B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-2001 (TrEMBLrel. 17, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Cellular tumor antigen p53 (Fragment).
TP53 OR P53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       D -> V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRINTS; PR00386; P53SUPPRESSR. ProDom; PD002681; P53; 1. PROSITE; P500348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL, X60020; CAA42635.1; -- HSSP; P04637; 10LG.
InterPro; IPR002117; P53.
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Les 18; Conservative
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             CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2

EXPRESSION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                              -!- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).
-!- DISEAGE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSPORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED IN MANY TYPES OF CANCER (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE P53 FAMILY.
-- SIMILARITY: AAH03596.1;
                                                                                                                                                                                                                                           Activator, Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein; Phosphorylation; Transcription regulation.
SEQUENCE 393 AA; 43686 MW; 5C701E458E016756 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE CELL DIVISION BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .;
0
                                                                                                                                                                                                                                                                                                     94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                    0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT; 393 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transformation suppressor (Fragment).
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Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
PRODOM; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
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es 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
                                                                                                                                                                HSSP; P04637; 10LG.
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                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606;
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SEQUENCE
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Q16535
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Plass M., Hartmann E., Mueller O., Kuhlmann J.;

Alfantification of a Sus scrofa gene encoding a protein with high
similarity to p53 from Homo sapiens, Bos taurus and Ovis aries.";
Submitted (JAN-1999) to the BMEL/GenBank/bDBJ databases.

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES
GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSTOCOGICAL
CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN
TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATE ARE INVOLVED IN
ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE RECLL DIVISION BY
CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE
ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
ACTIVATED GENES IS AN INHIBITOR OF STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2
EXPRESSION (BY SIMILARITY).
--- SUBGELLULAR LOCATION; NUCLEAR (BY SIMILARITY).
--- SUBGELLULAR LOCATION; NUCLEAR (BY SIMILARITY).
--- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF MANY TYPES OF CANCER.

MANY TYPES OF CANCER (BY SIMILARITY).
--- STANT AND TYPES OF CANCER (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Probon, P0002681, P33, 1.

Activator: Anti-oncogened Apoptosis; DNA-binding; Nuclear protein; Phosphorylation; Transcription regulation.

SEQUENCE 387 AA; 43475 MW; 3BB49A432B0053FA CRC64;
                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia! Eutheria; Cetartiodactyla; Suina; Suidae; Sus. NCBI TaxID=8923;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                  01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UJN-2002 (TrEMBLrel. 21, Last annotation update)
Cellular tumor antigen p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Search completed: February 12, 2003, 10:35:38
                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
TISSUE=URINARY BLADDER EPITHELIUM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
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Matches 17; Conservative
                                                                                                                                           Sus scrofa (Pig).
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Q9N252;
ПP
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neuropeptide - sea
gut pentapeptide -
bradykinin-potenti
Ig heavy chain CRD
Ig heavy chain CRD
T-cell receptor be
T-cell receptor be
Ig heavy chain CRD
alkanal monooxygen
endo-1,4-beta-xyla
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Met-enkephalin - b
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cocconase (EC 3.4.
T-call receptor be
serrawettin W2 - S
proctolin - Americ
peptidyl-dipeptida
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                          February 12, 2003, 10:49:54 ; Search time 15 Seconds (without alignments) 32.045 Million cell updates/sec
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GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 100 summaries
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A60803
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copper resistance flagellar protein R-phycocrythrin al zinc-binding prote proctolin - Atlant hypothetical prote

C41225 E42364 B22565 S55237 A60411 S53595

	ynaptosomal-asso	ell recept	-cell recept	7.5 kda stru	7.5K structu	/.sk structu	28.3A Structural pro	34.5K structural p	35K structural pro	surface protein te	phytosultokine alp	ribosomal protein	ribosomal protein	ribosomal protein	dnaZX-like protein	ı,	nhotosystem I 10 4	• .д	glycogen phosphory	peptidyl-dipeptida	COpper resistance	primase - Citrobac	- Citrob	20K protein - Rick	hypotherical prote	hypothetical prote	acid proteinase li	hypoxanthine phosp	Myosin iigne chain seminal plasma pro	. 0	T-ceil receptor be	T-cell receptor be	1-cell leceptor be 34.5K structural b	34.5K structural p	phosphoprotein, bo	spinal cord peptid	<b>"</b> (1)	ein - Es	tein	inding p	alpha-amylase - ri	hemoglobin, extrac	arconor denyarogen Ig heavy chain CRD	Ig heavy chain CRD	Ig kappa chain V-I	T-cell receptor be	ceptor	מָבָּ מַבָּ									
	2 D4482	5 2 PT0625	2 PT071	2 G4481	2 I4481	2 5446	2 A4481	2 B4481	2 D4481	2 86923	3 UTOB:	2 I3996	2 13996	2 I3996	2 I4046	2 E6027	2 PO066	2 \$6832	2 A6052	2 JN086	2 84122	2 I4070	2 I4069	2 B3183	2 T1490	2 T1491	2 B3798	2 A3/11	2 \$6288	2 B4482	2 PT059	2 PT059	2 H4481	2 F4481	2 \$1112	2 C2375	2 A4122	2 A3201	2 B6027	2 A3388	2 S5107	2 86572	2 PT026	2 PT029	2 JT052	2 C5328	2 PT061	2 PT051	2 PT052	2 PT060	2 PT068	2 PT060	2 PT067	2 PT066	2 PT065	2 PT065	2 PT065
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## ALIGNMENTS

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R.Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A.Title: Preferential utilization of specific immunoglobulin heavy chain diversity and j A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0308
                                                                                                                                                                                                                                                                                                                                                                                                                            C,Accession: G37196
R,Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
J. Protein Chem. 9, 227-227, 1990
A;Title: Primary structure and biological activity of bradykinin potentiating peptides A;Reference number: A37196; MUID:90351557; PMID:2386615
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PTO291
B;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
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C;Species: Homo sapiens (man)
C;Date: 30-Sep.1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0308
                                                                                                                                                                                                                                                                                                                                        bradykinin-potentiating peptide 7 - island jararaca
C;Species: Bothrops insularis (island jararaca)
C;Date: 14-Feb-1992 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994
                                                             Gaps
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A;Molecule type: protein
A;Molecule type: coro-
A;Molecule type: protein
A;Molecule type: doing acid
C;Keywords: pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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47.8%; Score 11; DB 2; Length 5; ilarity 100.0%; Pred. No. 2.8e+05; Conservative 0; Mismatches 0; Indels
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A, Experimental source: B lymphocyte
C, Keywords: heterotetramer; immunoglobulin
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Best Local Similarity
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A; Experimental source: gut
C; Comment: This peptide increased basal tone of the circular muscle of the esophagogastr
, and of the circular muscle of the gastro-intestinal junction.
                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-5 <SHI>
C;Comment: This peptide corresponds to the five carboxyl-terminal residues of cholecystd
                                                                                                                                                                                                                                                                           R;Shively, J.; Reeve Jr., J.R.; Bysselein, V.E.; Ben-Avram, C.; Vigna, S.R.; Walsh, J.H. An, J. Physiol. 252, G272-G275, 1987
A;Title: CCK-5: sequence analysis of a small cholecystokinin from canine brain and intes A;Reference number: A32516; MUID:87153871; PMID:3826354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brain Res. 442, 354-358, 1988
A;Title: Isolation of cGlu-Ser-Lu-Arg-Trp-NH-2, a novel neuropeptide from sea anemones.
A;Reference number: A60803; MUID:88222764; PMID:2897223
A;Accession: A60803.
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R;Ucesaka, T.; Ikeda, T.; Kubota, I.; Muneoka, Y.; Ando, M.
Biochem. Biophys. Res. Commun. 180, 828-832, 1991
A;Title: Structure and function of a pentapeptide isolated from the gut of the eel.
A;Reference number: JH0253; MUID:92062113; PMID:1953755
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N;Alternate names: CCK-5
C;Species: Canis lupus familiaris (dog)
C;Date: 18-Oct-1989 #sequence_revision 18-Oct-1989 #text_change 18-Aug-2000
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C;Species: Anguilla japonica (Japanese eel)
C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 11-Apr-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Species: Anthopleura elegantissima 30-Sep-1993 #text_change 07-May-1999 Ante: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999 Accession: Accession: Accession: C.J.P.
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A;Residues: 1-5 <GRA>
K;Rewindes: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;J/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;S/Modified site: amidated carboxyl end (Trp) #status experimental
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C;Keywords: amidated carboxyl end; neuropeptide
F;S/Modified site: amidated carboxyl end (Phe) #status experimental
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47.8%; Score 11; DB 2; Le
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
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A; Molecule type: protein
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A; Residues: 1-5 < YAM>

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alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi (fragment C;Species: Vibrio harveyi
C;Species: Vibrio harveyi
C;Species: Vibrio harveyi
C;Saccession: A44955
R;Paquatte, O.; Tu, S.C.
R;Paquatte, O.; Tu, S.C.
A;Title: Chemical modification and characterization of the alpha cysteine 106 at the Vibria A;Reference number: A44955; MUID:90175700; PMID:2626493
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Biochem. J. 316, 771-775, 1996
A;Title: Structural environment of an essential cysteine residue of xylanase from Chainia
A;Reference number: S70615; MUID:96265041; PMID:8670151
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A;Description: endohydrolyzation of beta-1,4-xylosidic linkages in xylans
A;Pathway: fermentation of hemicellulose into ethanol
C;Keywords: glycosidase; hydrolase
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Pred. No. 2.8e+05;
0; Mismatches 2; Indels
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A;Molecule type: protein
A;Residues: 1-5 <PAQ>
C;Keywords: FMN; luminescence; monooxygenase; oxidoreductase
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A,Molecule type: protein
A,Residues: 1-5 cRAD.
A,Experimental source: Chainia sp. strain NCL 82.5.1
A,Note: the source is designated as Chainia sp.
                                                                                                                                                                                                                                                                                                                            A;Molecule type: DNA
A;Residues: 1-5 <YAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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Best Local Similarity 50.0
Matches 2; Conservative
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reterence number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0640
A;Status: translation not shown
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C;Species: Mus musculus (house mouse)
C;Decies: Mus musculus (house mouse)
C;Decession: PT0580
R;Feeney, A.J.
J. Exp. Med. 115-124, 1991
A;Tible: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Wus musculus (house mouse)
C;Date: 17-Jul.1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0640; PT0685; PT0729
                                                                                                                                         Gaps
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A; Experimental source: newborn thymus, strain BALB/c, clone 120-1J
A; Experimental source: newborn thymus, strain BALB/c, clone 120-1J
A; Accession: PT0685
A; Status: translation not shown
A; Molecule type: DNA
A; Residues: 1-5 <FE2>
A; Residues: 1-5 <FE2>
A; Experimental source: day 18 fetal thymus, strain BALB/c, clone 154-1C
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A;Molecule type: DNA
A;Residues: 1-5 <FE3.
A;Experimental: ource: newborn thymus, strain BALB/c, clone 135-1AG
C;Keywords: T-cell receptor
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                                                                             Query Match 47.8%; Score 11; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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A,Residues: 1-5 <FEE>
A,Experimental source: day 19 fetal thymus, strain BALB/c
C,Keywords: T-cell receptor
A; Experimental source: B lymphocyte C; Keywords: heterotetramer; immunoglobulin
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Matches 1; Conservative
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C;Species: Achatina fulica (giant African snail)
C;Species: Achatina fulica (giant African snail)
C;Date: 23-Mar-1995 #sequence_revision 05-Apr-1995 #text_change 11-Jul-1997
C;Accession: A44692
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C;Species: Mytilus edulis (blue mussel)

C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000

C;Accession: B61445

R;Leung, M.K.; Stefano, G.B.

R;Leung, M.K.; St
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Met-enkephalin - blue mussel
C;Species: Mytilus edulis (blue mussel)
C;Decies: Mytilus edulis (blue mussel)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000
C;Accession: A61445
Expero, M.K.; Stefano, G.B.
Proc. Natl. Acad. Sci. U.S.A. 81, 955-958, 1984
A;Title: Isolation and identification of enkephalins in pedal ganglia of Mytilus edulis
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A;Residues: 1-5 <OHT>
C;Keywords: amidated carboxyl end; D-amino acid; neuropeptide
F;Z/Modified site: D-asparagine (Asn) #status experimental
F;S/Modified site: amidated carboxyl end (Val) #status experimental
                               A;Note: the authors translated the codon GAA for residue 3 as Gly C;Comment: The actin I gene contains no introns.
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A;Residues: 1-5 <LEU>
A;Experimental source: pedal ganglia
C;Keywords: neuropeptide; opioid peptide
A; Cross-references: GB:J03988
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C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Species: O3-Unr.1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C;Accession: B45525
R;Wesseling, J.G.; Snijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.; Schoenmak Mol. Biochem. Parasitol. 35, 16-176, 1989
A;Title: Stage-specific expression and genomic organization of the actin genes of the ma A;Reference number: A45525; MUID:89364996; PMID:2671721
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R; Maruyama, S.; Miyoshi, S.; Tanaka, H.
Agric. Biol. Chem. S3, 2763-2767, 1989
A; Tille: Angiotensin I-converting enzyme inhibitors derived from Ficus carica. A; Reference number: PQ0008
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.Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 02-Jul-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NiAlternate names: figus latex peptide 2
C;Species: Ficus carica (common fig)
C;Date: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 08-Dec-1995
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                                        Query Match 26.1%; Score 6; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0; Indele
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A,Residues: 1-5 <TSU>
A,Experimental source: leaf, chlorophyll
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A;Reference number: PS0206
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A;Residues: 1-5 <MAR>
A;Experimental source: latex
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A;Molecule type: DNA
A;Residues: 1-5 <WES>
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A; Reference number: PT0509; MUID:91277601; PMID:1711558
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Comp. Biochem. Physiol. C 88, 185-187, 1987
A;Title: Isolation and structure of a peptide isolated from the suboesophageal ganglion
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
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C;Species: Antheraea pernyi (Chinese oak silkmoth)
C;Species: Antheraea pernyi (Chinese oak silkmoth)
C;Accession: B61168
R;Kramer, K.J.; Felsted, R.L.; Law, J.H.
J. Biol. Chem. 248, 3021-3028, 1973
A;Fitle: Cocconase. V. Structural studies on an insect serine protease.
A;Feference number: A61168; MUID:73166540; PMID:4735570
                                                                                                                                                                                                                                                                                                                                                                       subesophageal ganglion pentapeptide - house cricket
C;Species: Acheta domesticus (house cricket)
C;Date: 07-Sep.1990 #sequence_revision 07-Sep-1990 #text_change 20-Jun-2000
C;Accession: JS0319
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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A;Molecule type: protein
A;Residues: 1-5 <KRA-
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-5/Product: cocoonase (fragment) #status experimental <MAT>
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A; Reference number: A61445; MUID:84144823; PMID:6583690 A; Accession: A61445
                                                                               A, Experimental source: pedal ganglia
C, Keywords: neuropeptide; opioid peptide
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                                                                                                                       Query Match
Best Local Similarity
                                        A; Molecule type: protein A; Residues: 1-5 < LEU>
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R; Feeney, A.J.
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C;Species: Serratia marcescens
C;Species: Serratia marcescens
C;bate: 10-Feb-1998 #sequence_revision 12-Feb-1998 #text_change 12-Feb-1998
C;Accession: A56728
R;Matsuyama, T.; Kaneda, K.; Nakagawa, Y.; Isa, K.; Hara-Hotta, H.; Yano, I.
Bacteriol. 114, 1769-1776, 1992
A;Title: A novel extracellular cyclic lipopeptide which promotes flagellum-dependent and A;Reference number: A58728; MUID:92193260; PMID:1548227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A/Molecule type: profein.
A/Residues: 1-5 <MAT>
A/Residues: 1-5 </MAT>
C/Commental Source: strain NS 25
C/Comment: A surfactant lipopeptide promoting flagellum-independent surface translocation C/Keywords: blocked amino end; blocked carboxyl end; D-amino acid; lipoprotein; unencodes F/Modified site: D-leucine (Leu) #stratus experimental
F/4/Modified site: D-phenylalanine (Phe) #status experimental
F/1/Cross-link: 3-hydroxydecanoyl amino end, ester carboxyl end (Leu-Ile) #status experimental
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C;Species: Periplaneta americana (American cockroach)
C;Date: 29-Jul-1981 #sequence_revision 29-Jul-1981 #text_change 23-Aug-1996
C;Accession: A01644
R;Starratt, A.N.; Brown, B.E.
Life Sci. 17, 1253-1256, 1975
A;Title: Structure of the pentapeptide proctolin, a proposed neurotransmitter in insects A;Reference number: A93048; MUID:76074708; PMID:576
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A; Residues: 1-5 < STA>
A; Residues: 1-5 < STA>
A; Note: the synthetic peptide had the same chromatographic, electrophoretic, and pharmack
B; O; Shea, M.; Adams, M.E.
Science 213, 567-569, 1981
A; Title: Pentapeptide (proctolin) associated with an identified neuron.
A; Title: Pentapeptide (proctolin) associated with an identified neuron.
A; Reference number: A94-866; WIDIS 81225865; PMID: 6113690
A; Contents: annotation; biological source
C; Comment: This peptide is found in the lateral white neurons, which occur (in the cockrismervate the striated hindgut muscles in insects and stimulate contraction of these musc
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
A Accession: PT0644
A;Status: translation not shown
A;Nolcule type: mRNA
A;Residues: 1-5 <FEE>
A;Residues: 1-C <FEE>
C;Keywords: T-cell receptor
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C;Species: Zea mays (maize)
C;Species: Zea mays (maize)
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 07-May-1999
C;Date: 27-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 07-May-1999
C;Accession: S55237
R;Robinson, K.; Jones, D.; Howell, S.; Soneji, Y.; Martin, S.; Aitken, A.
Biochem. J. 307, 267-272, 1995
A;Reference number: S55237; MUID:95234046; PMID:7717986
A;Accession: S55237; MUID:95234046; PMID:7717986
A;Accession: S55237; MUID:95234046; PMID:7717986
A;Reference number: S55237; MUID:95234046; PMID:7717986
A;Reference number: S55237; MUID:95234046; PMID:7717986
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R;Grome, J.R.; Tillinghast, B.K.; Townley, M.A.; Vetrovs, A.; Watson III, W.H.; Hunt, D peptides 11, 205-211, 1990
A;Title: Identification of proctolin in the central nervous system of the horseshoe crab A;Reference number: A60411; MUID:90287800; PMID:2356151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: protein
A;Residuee: 1-5 <05RO>
C;Comment: This neuropeptide stimulates cardiac output and hindgut motility in the horse
C;Keywords: neuropeptide
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C;Species: Limulus polyphemus (Atlantic horseshoe crab)
C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                      C;Species: Gastroclonium coulteri
C;Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C;Accession: B22565
R;Klotz, A.V.; Glazer, A.N.
J. Biol. Chem. 260, 4856-4863, 1985
A;Title: Chem. 260, 4856-4863, 1985
A;Reference number: A22565; MUID:85182601; PMID:3886644
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                        Length 5;
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                     17.4%; Score 4; DB 2; Ler 100.0%; Pred. No. 2.8e+05; tive 0; Mismatches 0;
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Best Local Similarity
Matches 1; Conservat
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A;Residues: 1-5 <KLO>
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A60411
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R. Matsumura, N.; Fujii, M.; Takeda, Y.; Shimizu, T.
Biosci. Blocechnol. Biochem. 57, 1743-11744, 1993
A. Title: Isolation and characterization of angiotensin I-converting enzyme inhibitory pe A; Reference number: JN0859; MUID: 94080036; PMID: 7764272
A; Reference number: JN0860
A; Molecule type: protein
A; Residues: 1-5 < MAT.
A; Residues: 1-5 < MAT.
A; Reperimental source: intestine
C; Comment: The carboxyl-terminus is essential for the protein's expression of angiotensic C; Superfamily: bradykinin-potentiating peptide
C; Superfamily: bradykinin-potentiating peptide
C; Keywords: angiotensin-converting enzyme inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C,Accession: C41225
R;Cha, J.S.; Cooksey, D.A.
R;Cha, J.S.; Cooksey, D.A.
Proc. Natl. Acad. Sci. U.S.A. 88, 8915-8919, 1991
A;Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer mem A;Reference number: A41225; MUID:92020961; PMID:1924351
A;Accession: C41225
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J. Bacteriol. 173, 3564-3572, 1991
A;Title: Salmonella typhimurium mutants defective in flagellar filament regrowth and A;Reference number: A42364; MUID:91258342; PMID:1646201
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C;Species: Sarda orientalis (striped bonito)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C.Species: Pseudomonas syringae pv. tomato
C.Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 24-Jun-1993
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C;Date: 24-Jul-1992 #sequence_revision 24-Jul-1992 #text_change 30-Sep-1993
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llarity 100.0%; Pred. No. 2.8e+05;
Conservative 0; Mismatches 0; Indels
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   0; Mismatches
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A,Residues: 1-5 <VOG>
A,Cross-references: GB:M62408
   Conservative
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Best Local Similarity
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C,Accession: PT0625
R;Feeney, A.J.
J. Exp. Med. 114, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                 T-cell receptor beta chain V-D-J region (120-25) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
                                                                                                                                           17.4%; Score 4; DB 2; Length 5;
100.0%; Pred. No. 2.8e+05;
cive 0; Mismatches 0; Indels
                                      A;Residues: 15 <FEE>
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-5 <FBE>
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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                                                                                                                          Query Match
Best Local Similarity 100.0
Matches 1; Conservative
             A; Molecule type: mRNA
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                                                                                                                                                                                                                                                          hypothetical protein (upstream of transcription factor, CCAAT-binding) - chicken C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Date: 15-Jul-1995 #sequence_revision 01-Sep-1995 #text_change 07-May-1999
C;Accession: S55595
R;Calkhoven, C.F.; Bouwman, P.R.J.; Snippe, L.; Ab, G.
Nucleic Acids Res. 22, 5540-5547, 1994
A;Title: Translation start site multiplicity of the CCAAT/enhancer binding protein alpha A;Reference number: S53595; MUID:95140613; PMID:7838705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synaptosomal-associated protein SNAP-25 peptide 2 - rabbit (fragment)
N'Alternate names: superprotein peptide 2
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996
C;Accession: D44823
R;Loewy, A.; Liu, W.S.; Baitinger, C.; Willard, M.B.
J. Neurosci. 11, 3412-3421, 1991
A;Reference number: A44823; MUID:92044785; PMID:1941090
A;Reference number: A44823; MUID:92044785; PMID:1941090
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C;Species: Mus musculus (house mouse)
C;Decies: Mus musculus (house mouse)
C;Decies: JJ-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0624
A;Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Feeney, A.J.
A;Fitele: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0624
A;Status: translation not shown
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indel
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A.Residues: 1-5 <LOE>
A.Experimental source: visual tissue
A.Note: sequence extracted from NCBI backbone (NCBIP:64250)
C.Keywords: membrane trafficking
Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0;
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A;Molecule type: DNA
A;Residues: 1-5 <CAL>
A;Cross-references: BMBL:X66844
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J. Exp. Med. 174, 115-124, 1991
A;Title: Unctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Leuconostoc oenos phage P32
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: G44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indel
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17.4%; Score 4; DB 2; Les
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
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A;Molecule type: DNA
A;Residues: 1-5 <FEE
C;Keywords: T-cell receptor
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C;Species: Leuconostoc oenos phage P37
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: 144817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUD: 92085033; PMID:1748868
A;Molecule type: protein
A;Residues: 1-5 <ARE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27.5K structural protein - Leuconostoc oenos phage P54 (fragment)
C;Species: Leuconostoc oenos phage P54
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                         A; Note: sequence extracted from NCBI backbone (NCBIP:70333)
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
d. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: E44817
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: G44817
A;Molecule type: protein
A;Residues: 1-5 <ARE>
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C;Species: Leuconostoc oenos phage PAt5-12
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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C,Species: Leuconostoc oenos phage PZt11-15
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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C;Species: Leuconostoc oenos phage PZt11-15
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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A;Residues: 1-5 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70342)
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: G44817
A;Molecule type: protein
A;Residues: 1-5 < ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70341)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Note: sequence extracted from NCBI backbone (NCBIP:70343)
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R; Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A; Title: Lysogeny in Leuconostoc oenos.
A; Reference number: A44817; MUID:92085033; PMID:1748868
A; Accession: B44817
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Tille: Lysogeny in Leuconostoc Oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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A;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J; Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
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Libosomal protein S4 - Bacillus circulans (fragment)
C,Species: Bacillus circulans
C,Species: Bacillus circulans
C,Species: Bacillus circulans
C,Species: 19-Jul-1996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
C,Accession: 13964
R;Grundy, F.J.; Henkin, T.M.
A,Title: Characterization of the Bacillus subtilis rpsD regulatory target site.
A,Accession: 13964
A,Accession: 13964
A,Accession: 13964
A,Accession: 13964
A,Reterence number: 13963; MUID:93015735; PMID:1400226
A,Accession: 13964
A,Accession: 15964
A,Accession: 15964
A,Accession: IS964
A,Residues: 1-5 <-RES>
A,Accession: IS964
A,Residues: 1-5 <-RES>
A,Accession: IS964
A,Residues: 1-5 <-RES>
A,Accession: IS964
A,Residues: ISON A,Accession: IS964
A,Accession: IS964
A,Residues: ISON A,Accession: IS964
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C;Species: Bacillus licheniformis
C;Species: 19-Jul-1996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
C;Accession: 13996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
B;Grundy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 6763-6770, 1992
A;Title: Characterization of the Bacillus subtilis rpsD regulatory target site.
A;Reference number: 139963; MUID:93015735; PMID:1400226
A;Accession: 139966
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13.0%; Score 3; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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13.0%; Score 3; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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C;Species: Bacillus megaterium
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A;Residues: 1-5 <RES>
A;Cross_references: GB:M99043; NID:g143475
                                                    A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-5 <BRA>
A;Cross-references: GB:M63747
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C;Species: Staphylothermus marinus (fragment)
C;Date: 04-Dec-1997 #sequence_revision 04-Dec-1997 #text_change 17-Mar-1999
C;Date: 04-Dec-1997 #sequence_revision 04-Dec-1997 #text_change 17-Mar-1999
C;Accession: 869237
Mol. Biol. 245, 385-401, 1995
J. Mol. Biol. 245, 385-401, 1995
J. Mol. Biol. 245, 385-401, 1995
A;Title: Tetrabrachion: a filamentous archaebacterial surface protein assembly of unusua A;Reference number: 869237; MUID:95139068; PMID:7837271
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C;Date: 11-Sep-1992 #sequence_revision 11-Sep-1992 #text_change 23-Mar-1993
C;Accession: B37325
R;Braaten, B.A.; Blyn, L.B.; Skinner, B.S.; Low, D.A.
J. Bacteriol. 173, 1789-1800, 1991
A;Title: Evidence for a methylation-blocking factor (mbf) locus involved in pap pilus ex
A;Reference number: A37325; MUID:91154136; PMID:1671857
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Proc. Natl. Acad. Sci. U.S.A. 93, 7623-7627, 1996
A;Title: Phytosulfokine, sulfated peptides that induce the proliferation of single mesop
A;Reference number: JT0870
A;Accession: JT0870
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C;Species: Asparagus officinalis (garden asparagus)
C;Date: 02-Apr-2002 #sequence_revision 02-Apr-2002 #text_change 19-Apr-2002
C;Accession: JT0870
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C;Species: Escherichia coli
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                                                                                                                                                                                                                                   / Match 17.4%; Score 4; DB 2; Length 5; Local Similarity 100.0%; Pred. No. 2.8e+05; les 1; Conservative 0; Mismatches 0; Indel
                                                                         A;Molecule type: protein
A;Residues: 1-5 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70340)
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A;Residues: 1-5 <PET>
A;Experimental source: strain F1, DSM 3639
C;Keywords: cell wall; glycoprotein; heat-stable protein
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: D44817
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Matches
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NyAlernate names: Abcp-B
C;Species: Ascidia ceratodes
C;Date: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 11-Jun-1999
C;Accession: S68326
R;Taylor, S.W.; Ross, M.M.; Waite, J.H.
Rxch. Biochem. Biophys. 324, 228-240, 1995
A;Title: Novel 3,4-di- and 3,4,5-trihydroxyphenylalanine-containing polypeptides from th
A;Reference number: S68325; MUID:96132650; PMID:8554314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Nicotiana tabacum (common tobacco)
C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 17-Mar-1999
C;Date: 19-May-1994 #sequence_revision 19-May-1994 #text_change 17-Mar-1999
C;Accession: PQ0689
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A;Molecule type: protein
A;Residues: 1-5 <TAY>. 5 <TAY-7
F;2/Modified site: 3',4',5'-trihydroxyphenylalanine (Tyr) #status experimental
F;4/Modified site: 3',4'-dihydroxyphenylalanine (Tyr) #status experimental
                                                                                   P.phycoerythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) (Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Anar-1998 #sequence_revision 07-Mar-1998 #text_change 23-Mar-1993 C,Accession: F22565 A.N. i Glazer, A.N. i Glazer, A.N. i Glazer, A.N. i Glazer, A.N. i A.Y. i A
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A;Residues: 1-5 <080>
C;Keywords: chloroplast; photosynthesis; photosystem I; thylakoid
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R;Nagai, S.; Wiker, H.G.; Harboe, M.; Kinomoto, M.
Infect. Immun. 59, 372-382, 1991
A;Title: Isolation and partial characterization of major protein antigens in the culture A;Reference number: A60274; MUID:91099989; PMID:1898899
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C;Species: Bacillus subtilis
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 21-Jul-2000
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 21-Jul-2000
C;Accession: 140469
R;Struck, J.C.; Hartmann, R.K.; Toschka, H.Y.; Erdmann, V.A.
Mol. Gen. Genet. 215, 478-482, 1989
A;Aritle: Transcription and processing of Bacillus subtilis small cytoplasmic RNA.
A;Reference number: 140469; MUID:89218958; PMID:2468993
A;Accession: 140469
A;Accession: preliminary; translated from GB/EMBL/DDBJ
A;Residues: 1-5 <RES>
A;Residues: 1-5 <RES>
A;Cross-references: EMBL:X14796; NID:940130; PIDN:CAA32902.1; PID:94376204
A;Start codon: GTG
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                               C,Accession: I39965
R,Grundy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 673-6770, 1992
A,Title: Characterization of the Bacillus subtilis rpsD regulatory target site.
A,Reference number: 139963; MUID:93015735; PMID:1400226
A,Accession: 139963; MUID:93015735; PMID:1400226
A,Status: preliminary; translated from GB/EMBL/DDBJ
A,Rolecule type: DNA
A,Residues: 1-5 <RES>
A,Cossereferences: GB:M99042; NID:g143473
A,Genetics:
A,Genetics:
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C;Species: Mycobacterium tuberculosis
C;Date: 11-Dec-1992 #sequence_revision 11-Dec-1992 #text_change 30-Sep-1993
C;Date: 19-Jul-1996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
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Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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primase - Citrobacter diversus (fragment)
C;Species: Citrobacter diversus
C;Species: Citrobacter diversus
C;Date: 16-Aug-1996 #sequence_revision 16-Aug-1996 #text_change 16-Aug-1996
C;Accession: 140702
R;Versalovic, J.; Lupski, J.R.
Mol. Microbiol. 8, 343-355, 1993
A;Title: Conservation and evolution of the rpsU-dnaG-rpoD macromolecular synthesis (MMS)
A;Reference number: 140702; MUID:93302510; PMID:8316085
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C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Datesion: 146698
B;Shiuan, D.; Campbell, A.
Gene 67, 203-211, 1988
A;Title: Transcriptional regulation and gene arrangement of Escherichia coli, Citrobacte A;Reference number: 140697; MUID:89006280; PMID:2971595
A;Reference number: 140698
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
                                                                                                                                                                                                                             C;Species: Xanthomonas sp. C. C;Species: Xanthomonas sp. C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 08-Oct-1999 C;Accession: 870154 #sequence_revision 13-Mar-1997 #text_change 08-Oct-1999 C;Accession: 870161; G:Y:, Mindlin, S:Z:; Bass, I.A.; Yurieva, O.V.; Minakhina, S:V.; Nikiforov, Mol. Microbiol. 17, 1189-1200, 1995 A;Title: Four genes, two ends, and a res region are involved in transposition of Th5053: A;Reference number: 870140; MUID:96130850; PMID:8594337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-5 < KHO>
A; Cross-references: EMBL:L40585; NID:g710572; PIDN:AAA98329.1; PID:g735909
A; Note: the nucleotide sequence was submitted to the EMBL Data Library, May 1995
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Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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A;Molecule type: DNA
A;Residues: 1-5 <RES>
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                             glycogen phosphorylase (EC 2.4.1.1), muscle - mullet (Liza ramada) (fragment)
N.Alternate names: glycogen phosphorylase b
C;Species: Liza ramada
C;Species: Liza ramada
C;Species: Liza ramada
C;Accession: A60521
R;Bonamusa, L.; Baanante, I.V.
C;Accession: A60521
R;Bonamusa, L.; Baanante, I.V.
A;Mille: Purification and characterization of glycogen phosphorylase B from skeletal muscle A;Reference number: A60521; MUID:90227907; PMID:2109669
                                                                                                                                                                                                                                                                                                                                    A;Molecule type: protein
A;Medidues: 1-5 - 8DN>
A;Residues: 1-5 - 8DN>
C;Superfamily: phosphorylase
C;Superfamily: phosphorylase
C;Keywords: glycosyltransferase; hexosyltransferase; phosphoprotein
F;3/Binding site: phosphate (Ser) (covalent) (by phosphorylase b kinase) #status experin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       peptidy1-dipeptidase A inhibitory peptide C112 - striped bonito
C;Species: Sarda orientalis (striped bonito)
C;Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
C;Accession: JN0862
R;Matsumura, N.; Fujii, M.; Takeda, Y.; Shimizu, T.
Biosci. Biotechnol. Biochem. S7, 1743-1194, 1993
A;Title: Isolation and characterization of angiotensin I-converting enzyme inhibitory pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A) Experimental source: intestine
C; Comment: The amino terminal tripeptide of this protein inhibits angiotensin I-converti
C; Superfamily: bradykinin-potentiating peptide
C; Keywords: angiotensin-converting enzyme inhibitor
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Cispecies: Pseudomonas syringae pv. tomato
Cispecies: Pseudomonas syringae pv. tomato
Cibate: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 24-Jun-1993
Ciscession: B41225
R;Cha, J.S.; Cooksey, D.A.
R;Cha, J.S.; Cooksey, D.A.
R;Cha, J.S.; Cooksey, D.A.
A;Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer mem A;Reference number: A41225; MUID:92020961; PMID:1924351
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Best Local Similarity 0.0%
Matches 0; Conservative
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A,Molecule type: protein
A,Residues: 1-5 <MAT>
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A;Molecule type: protein
A;Residues: 1-5 <CHA>
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RESULT 52
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hypothetical protein - parsley
C;Species: Petroselinum crispum (parsley)
C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 21-Jul-2000
C;Accession: T14910
R;Kircher, S.; Ledger, S.; Hayashi, H.; Weisshaar, B.; Schafer, E.; Frohnmeyer, H.
Mol. Gen. Cenet. 257, 595-605, 1998
A;Title: CPRF4a, a novel plant bZIP protein of the CPRF family: comparative analysis of
A;Reference number: Z18261; MUID:98265918; PMID:9604882
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R;Murakami-Murofushi, K.; Takahashi, T.; Minowa, Y.; Iino, S.; Takeuchi, T.; Kitagaki-Og
N; Biol. Chem. 265, 19898-19903, 1990
A;Title: Purification and characterization of a novel intracellular acid proteinase from A;Reference number: A37988; MUID:91060608; PMID:2246266
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C'Date: 28 Mar-1991 #sequence_revision 28-Mar-1991 #text_change 23-Jun-1993
C'Accession: A37114
R'Yuan, L.; Craig, S.P.; McKerrow, J.H.; Wang, C.C.
A' Biol. Chem. 265, 13528-13532, 1990
A'Title: The hypoxanthine phosphoribosyltransferase of Schistosoma mansoni. Furt
A;Reference number: A37114; MUID:90337955; PMID:2199439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary, translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-5 <KINA
A;Cross-references: EMBL:Y10810; NID:93336904; PIDN:CAA71769.1; PID:93336905
A;Experimental source: ssp. Hamburger Schnitt
                                      A;Cross-references: EMBL:Y10809; NID:g3336901; PIDN:CAA71767.1; PID:g3336902 A;Experimental source: Hamburger Schnitt
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                                                                                                                                                           Query Match 8.7%; Score 2; DB 2; Length 5; Best Local Similarity 0.0%; Pred. No. 2.8e+05; Matches 0; Conservative 1; Mismatches 0; Indels
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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A;Molecule type: protein
A;Residues: 1-5 <MUR>
A; Residues: 1-5 < KIR>
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c; Species: Vicia sativa (spring vetch, tare)
C; Species: Vicia sativa (spring vetch, tare)
C; Species: Vicia sativa (spring vetch, tare)
C; Date: 16-Uul-1999 #sequence_revision 16-Uul-1999 #text_change 16-Uul-1999
C; Accession: T10994 #sequence_revision 16-Uul-1999 #text_change 16-Uul-1999
R; Christiansen, A.; Hansen, A.C.; Vijn, I.; Pallisgard, N.; Larsen, K.; Yang, W.C.; Biss submitted to the EMBL Data Library, December 1995
A; Description: A novel type of DNA binding protein interacts with a conserved sequence in A; Reference number: Z1728
A; Accession: T1054
A; Accession: T1055
A; Molecule type: mRNA
A; Residues: 1-5 < CHR>
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Cydicesion: B31836
R;Anderson, B.E.; Baumstark, B.R.; Bellini, W.J.
J. Bacteriol. 170, 4493-4500, 1988
A;Title: Expression of the gene encoding the 17-kilodalton antigen from Rickettsia ricke
A;Reference number: A91885; MUID:89008059; PMID:3139629
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C;Species: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 21-Jul-2000
C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 21-Jul-2000
C;Accession: T14908
R;Kircher. S;; Ledger, S; Hayashi, H.; Weisshaar, B.; Schafer, E.; Frohnmeyer, H. Mol. Gen. Gent. 257, 595-605, 1998
A;Title: ;CPRF4a, a novel plant bZIP protein of the CPRF family: comparative analysis of A;Reference number: Z18261; MUID:98265918; PMID:9604882
A;Accession: T14908
A;Accession: T14908
A;Accession: T14908
A;Accession: MUID: MUID:
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                                                                                               Length 5;
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                                                                                  8.7%; Score 2; DB 2; L. Local Similarity 0.0%; Pred. No. 2.8e+05; les 0; Conservative 1; Mismatches 0
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        A; Cross-references: GB: M21922; NID: g144434
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C;Accession: B44823

A; Liu, W.S.; Baitinger, C.; Willard, M.B.

J. Neurosci. 11, 3412-3421, 1991

A;Title: The major 35S-methionine-labeled rapidly transported protein (superprotein) is

A;Reference number: A44823; MUID:92044785; PMID:1941090

A;Status: preliminary
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C;Species: Mus musculus (house mouse)
C;Species: Mus musculus (house mouse)
C;Accession: P710590
R;Feeney, A.J.
J. Exp. Med. 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: P70509; MUID:91277601; PMID:1711558
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J. Exp. Med. 174, 115-124, 1991
J. Exp. Med. 174, 115-124, 1991
A.JTILE: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A.Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0597
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                                                                                                                                                                                       A,Molecule type: protein
A,Residues: 1-5 <LOB>
A,Experimental source: visual tissue
A,Note: sequence extracted from NCBI backbone (NCBIP:64255)
C,Keywords: membrane trafficking
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C;Keywords: T-cell receptor
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A;Residues: 1-5 <FEE>
A;Experimental source: newborn thymus, strain BALB/c
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A,Molecule type: mRNA
A,Residues: 1-5 <FEE>
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PT0597
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C;Species: Gallus gallus (chicken)

C;Species: Gallus gallus (chicken)

C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 28-Feb-1997

C;Accession: 150385

R;Shen, R.; Goswami, S.K.; Mascareno, E.; Kumar, A.; Siddiqui, M.A.Q.

R;Shen, R.; Goswami, S.K.; Mascareno, E.; Kumar, A.; Siddiqui, M.A.Q.

A;Title: Tissue-specific transcription of the cardiac myosin light-chain 2 gene is regul

A;Reference number: 150385; MUID:91141519; PMID:1996116

A;Accession: 150385

A;Status: preliminary; translated from GB/EMBL/DDBJ
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FEBS Lett. 382, 15-17, 1996
A;Title: Crystallization and preliminary X-ray diffraction analysis of boar seminal plas A;Reference number: S62882; MUID:96196555; PMID:8612739
A;Accession: S62883
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A;Residues: 1-5 <ROM>
C;Complex: heterodimer; seminal plasma protein I and seminal plasma protein
C;Keywords: glycoprotein; heterodimer; semen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     seminal plasma protein II - pig (fragment)
C;Species: Sus scrofa domestica (domestic pig)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
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N;Alternate names: superprotein peptide 10A
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996
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                         A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-5 <YUA>
C;Keywords: glycosyltransferase; pentosyltransferase
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C;Genetics:
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Best Local Similarity 0.0%
Matches 0; Conservative
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     A; Accession: A37114
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phosphoprotein, bone - chicken (fragment)
C;Species; Gallus gallus (chicken)
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
C;Accession: S11127; S11128
R;Mikuni-Takagaki, Y.; Glimcher, M.J.
Biochem, J. 268, 588-591, 1990
A;Title: Post-translational processing of chicken bone phosphoproteins. Identification A;Reference number: S11127; MUID:90303246; PMID:2363696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Spinal cord peptide SCP-6 - pig
Cispecies: Sus scroff adomestica (domestic pig)
Cibate: 28-89-1987 #sequence_revision 28-8ep-1987 #text_change 18-Aug-2000
CiAccession: C23751
Rihsi, K.L.; Chen, R.L.; Chen, Z.G.; Zhang, H.L.; Lu, Y.A.; Guo, S.Y.; Wu, S.X.; Tsou,
Arch. Biochem. Biophys. 240, 178-183, 1985
A;Reference number: A23751; MUID:85250425; PMID:4015098
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C;Species: Mus musculus (house mouse)
C;Accession: A26830
R;Reichelt, K.; Elgjo, K.; Edminson, P.D.
B;chem. Biophys. Res. Commun. 146, 1493-1501, 1987
A;Title: Isolation and structure of an epidermal mitosis inhibiting pentapeptide.
A;Reference number: A26830; MUID:87298602; PMID:3619940
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4.3%; Score 1; DB 2; L
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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A;Molecule type: protein
A;Residues: 'X', 2-5 <MIK2>
C;Keywords: phosphoprotein
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A;Molecule type: protein
A;Residues: 1-5 <MIKl>
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A;Molecule type: protein
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      34.5K structural protein - Leuconostoc oenos phage P32 (fragment)
C;Species: Leuconostoc oenos phage P32
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: H44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gan. Microbiol. 137, 2135-2139, 1991
A;Title: Lysegeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Molecule type: protein
A;Residues: 1-5 <ARE>
                                                 T-cell receptor beta chain V-D-J region (141-100) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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C;Species: Leuconostoc oenos phage P54
C;Date: 31-Mar_1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                  A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-5 <FEE.
A;Experimental source: day 19 fetal thymus, strain BALB/c
C;Keywords: T-cell receptor
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: F44817
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Copper resistance protein A - Pseudomonas syringae pv. tomato (fragment)
C,Species: Pseudomonas syringae pv. tomato
C,Species: Pseudomonas syringae pv. tomato
C,Species: 19-3un-1992 #sequence_revision 19-3un-1992 #text_change 24-3un-1993
C,Accession 441225
R,Cha, J.S.; Cooksey, D.A.
Proc. Natl. Acad. Sci. U.S.A. 88, 8915-8919, 1991
A,Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer mem A,Reference number: A41225; MUID:92020961; PMID:1924351
A,Accession: A41225
A,Accession: A41225
A,Reterence number: A41225
A,Reterence num
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Best Logal Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 0; Mismatches 1; Indels
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

February 12, 2003, 10:46:33; Search time 11 Seconds (without alignments) 18.853 Million cell updates/sec

US-09-403-440A-1 Title: Perfect score:

1 FXXLW 5 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 segs, 41476328 residues Searched:

20 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 5 Maximum DB seq length: 5

Maximum Match 100% Listing first 100 summaries Post-processing: Minimum Match 0%

SwissProt\_40:\* Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result		* Ouerv					
No.	Score	Match	Match Length DB	B0	ΩI	Description	tion
1	11	47.8	S	7	BPP7 BOTIN	P30425	bothrops in
7	11	47.8	S	П	UF01 MOUSE	P38639	mus musculu
٣	9	26.1	ιΩ	٦	AL14 CARMA	P81817	carcinus ma
4	9	26.1	S	н	EIO3 LITRI	J P82099	litoria rub
Ŋ	9	26.1	Ŋ	Н	EIO4 LITRU	J P82100	litoria rub
9	9	26.1	S	П	FARP ARTTE	P41853	artioposthi
7	9	26.1	S	-	PAP2 PARM	A P81864	pardachirus
80	9	26.1	Ŋ	-	RE11 LITR	J P82070	litoria rub
6	9	26.1	S	-	RE21 LITR	P82071	litoria rub
10	9	26.1	S	-	RE31 LITE	P82072	litoria rub
11	y	26.1	Ŋ	H	RE32 LITR	U P82073 1	litoria rub
12	ø	26.1	Ŋ	Н	SUGA_ACHDO	P19991	acheta dome
13	9	26.1	S	Н	TPIS CANF.	P54714	canis famil
14	9	26.1	Ŋ	Н	UC22 MAIZE	P80628	zea mays (m
15	4	17.4	S	٦	PRCT PERAM	M P01373	periplaneta
16	4	17.4	2	٦	PSK DAUCA	P58261	daucus caro
17	7	8.7	2	Н	BIOA CITFR	P13071	citrobacter
18	7	8.7	Z,	-	BIOB CITFR	P12997	citrobacter
19	0	0.0	5	Н	TRM3 ECOL	I P13973	escherichia
20	0	0.0	Ŋ	ч	UXA4 CHLTR	P38005	chlamydia t

## ALIGNMENTS

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01-APR-1993 (Rel. 25, Created)
01-FBB-1994 (Rel. 28, Last sequence update)
01-FBB-1994 (Rel. 28, Last annotation update)
Bradykinin-potentiating peptide 55,2 (5A) (Angiotensin-converting
                             5 AA.
                             STANDARD;
                             BOTIN
                           BPP7 BOT
P30425;
RESULT 1
BPP7 BOTIN
                             55 PT PT PE
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                                                                                                                                                                           Cintra A.C.O., Vieira C.A., Giglio J.R.;

"Primary structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";

J. Protein Chem. 9:221-227(1990).

-!- FUNCTION: THIS PEPTIDE BOTH INHIBITS THE ACTIVITY OF THE ANOIOTENSIN-CONVERTING BNZYME AND ENHANCES THE ACTION OF BRADYKININ BY INHIBITING THE KINASES THAT INACTIVATE IT.

IT ACTS AS AN INDIRECT HYPOTENSIVE AGENT.

PIR; G37196; G37196.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
Bothrops insularis (Island jararaca) (Queimada jararaca).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.1e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                               1 1 PYRROLIDONE CARBOXYLIC ACID
5 AA; 629 MW; 776DC37326B00000 CRC64;
                                                                                                                                           TISSUE=Venom;
MEDLINE=90351557; PubMed=2386615;
                                                                                                                                                                                                                                                                                                                                              Hypotensive agent; Venom. MOD RES 1 1
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 01-0CT-1994 (Rel. 30, Created) 01-0CT-1994 (Rel. 30, Last sequence update) 01-0CT-1995 (Rel. 31, Last annotation update) Unknown protein from 2D-page of fibroblasts (P19) (Fragment). 5 AA. STANDARD; Mus musculus (Mouse) NCBI\_TaxID=10090; UF01 MOUSE P38639; UF01\_MOUSE

Electrophoresis 15:735-745(1994).
-!- MISCELLANBOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 6.6, ITS MW IS: 19 kDa.
NON\_TER TISSUE=Fibroblast;
MEDLINE=55009907; PubMed=7523108;
Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;
"Separation and sequencing of familiar and novel murine proteins using preparative two-dimensional gel electrophoresis."; 5 AA; 717 MW; 7364087043100000 CRC64; SEQUENCE SEQUENCE 

ò Gaps ; 0 Indels 47.8%; Score 11; DB 1; Length 5; 100.0%; Pred. No. 1.1e+05; 100.0%; Pred. nc. Conservative Local Similarity 5 14 5 N. Matches RESULT g ઠ

Query Match

(Rel. 39, Created) (Rel. 39, Last sequence update) (Rel. 39, Last annotation update) STANDARD; P81817; 30-MAY-2000 ( 30-MAY-2000 ( 30-MAY-2000 ( AL14 CARMA A114 CARMA
ID A114 CP
AC P818I7;
DT 30-MAYDT 30-MAYDT 30-MAYDT 30-MAY-

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Litoria rubella (Desert tree frog).
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P81864;
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Carcinus maenas (Common shore crab) (Green crab).
Bukaryota, Metazoa, Arthropoda; Mandibulata; Pancrustacea; Crustacea;
Malacostraca; Eumalacostraca; Eucarida; Decapoda; Pleocyemata;
Brachyura; Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBI_TaxID=6759;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C.; "Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Litoria rubella (Desert tree frog).
Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Amphibia, Batrachia, Anura; Neobatrachia, Bufonoidea, Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
                                                                                                                                                                                                                                                                       "Isolation and identification of multiple neuropeptides of the
                                                                                                                                                                                                                                                                                             allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250:77-734 (1997).
-1- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-1- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
5 S AMIDATION (POTEWILL).
SEQÜENCE 5 AA; 586 MW; 672879D5AB300000 CRC64;
                                                                                                                                                                                                               Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P., Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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5 AA; 630 MW; 668761F2C9A00000 CRC64;
                                                                                                                                                                             TISSUE-Cerebral ganglion, and Thoracic ganglion, MEDLINE-98121193; PubMed-9461295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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-!- SUBCELLULAR LOCATION: SECRETED.
Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
nes 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI TaxID=104895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E103 LITRU
P82099;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E104 LITRU
P82100;
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EI04_LITRU
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Matches
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Platyhelminthes; Turbellaria; Seriata; Tricladida;
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                                                                                                                                                                                         TISSUE=Skin secretion;
Wabhitz P.A., Bowle J.H., Tyler M.J., Wallace J.C.;
Wabhitz From the skin glands of the Australian buzzing tree frog
Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Pardachirus marmoratus (Red sea moses sole).
Pardachirus marmoratus (Red sea moses sole).
Pardachirus marmoratus (Red sea moses sole).
Actinopterygii, Neopterygii, Teleostei; Buteleostei; Neoteleostei; Acanthomorpha; Acanthopterygii, Percomorpha; Pelantonectiformes; Soleoidei; Soleidae; Pardachirus.
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=94211927; PubMed=7909164;
Maule A.G., Shaw C., Halton D.W., Curry W.J., Thim L.;
"RYIRFamide: a turbellarian FMRFamide-related peptide (FaRP).";
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-!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match 26.1%; Score 6; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 1.1e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE 5 AA; 616 MW; 61F2D1A059A00000 CRC64;
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SEQUENCE 5 AA; 754 MW; 69D4004B44600000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                               AMIDATION
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Terricola, Geoplanidae, Arthurdendyus.
                                                                                                                                                                                                                                                                                                   Aust. J. Chem. 52:639-645 (1999).

-I- SUBCELLULAR LOCATION: SECRETED.
Amphibian skin; Amidation.

AMIDAT
AMD PR. 5
AMIDAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Neuropeptide; Amidation.
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                                                                                         NCBI TaxID=104895;
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15-JUN-2002 (Rel. 41, Last annotation update)
                Rubellidin 2.1.
Litoria rubella (Desert tree frog)
                                                                           NCBI_TaxID=104895;
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                                                                                                                                                                                                                                                 Amphibian skin.
SEQUENCE 5 AA
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                                                                                                                                                                                                               ACTIVITY
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                                                                                                                                                                                                                                                                                                                                                                                                          RE31 LITRU
ID RE31 LITRU
AC P82072;
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RE32_LITRU
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                                 TISSUE=Skin secretion;
MEDLINE=1057369; PubMed=3782138;
Lazarovici P., Primor N., Loew L.M.;
Lazarovici P., Primor N., Loew L.M.;
Purification and pore-forming activity of two hydrophobic polypeptides from the secretion of the Red sea moses sole (Pardachirus marmoratus).'.

Biol. Chem. 261:16704-16713(1986).

-!- FUNCTION: EXHIBITS UNUSUAL SHARK REPELLENT AND SURFACTANT PROPERTIES. FORMS VOLTAGE-DEPENDENT, ION-PERMEABLE CHANNELS IN MEMBRANES. AT HIGH CONCENTRATION CAUSES CELL MEMBRANE LYSIS.
-!- SUBGNAIT: MONOWER. IN ADUEDOUS SOLUTION EXISTS AS A TETRAMER.
-!- SUBCELULAR LOCATION: Secreted.
-!- SIMILARITY: BELONGS TO THE PARDAXIN FAMILY.
                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The structure of new peptides from the Australin red tree frog Litoria rubella. the skin peptide profile as a probe for the study of evolutionary trends of amphibians.",
Aust. J. Chem. 49:955-963 (1996).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                       Rubellidin 1.1.
Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Amphibia; Batrachia, Anvra; Neobatrachia; Bufonoidea; Hylidae, Pelodryadinae; Litoria.
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0
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Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
Tyler M.J., Wallace J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -1- TISSUE SPECIFICITY: SECRETED BY THE SKIN DORSAL GLANDS
-1- MASS SPECTROMETRY: MW=598; METHOD=FAB.
                                                                                                                                                                                                                                           Match 26.1%; Score 6; DB 1; Length 5; Local Similarity 100.0%; Pred. No. 1.18+05; les 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
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                                                                                                                                                                                                         5 AA; 614 MW; 7769C9C9C8100000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 AA; 598 MW; 6DD9C9CAB2A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                  15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                 5 AA.
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(Rel. 41, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE, AND MASS SPECTROMETRY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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WCBI TaxID=31087;
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P82071;
15-JUN-2002
15-JUN-2002
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P82070;
                                                                                                                                                                                                         NON TER
SEQUENCE
                         SEQUENCE
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                                                                                                                                                                                                                                            Query Match
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RE21 LITRU
                                                                                                                                                                                                                                                                   Matches
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TISSUE-Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Trjet M.J., Waller G. J.C.,
"The structure of new peptides from the Australin red tree frog
Litoria rubella'. the skin peptide profile as a probe for the study
of evolutionary trends of amphibians.";
Aust. J. Chem. 49:955-963(1996).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Aust. J. Chem. 49:955-963(1996).

-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOFIC
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia; Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae;
Pelodryadinae, Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Litoria rubella (Desert tree frog).
Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria
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Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
Tyler M.J., Wallace J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 26.1%; Score 6; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 1.1e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26.1%; Score 6; DB 1; Length 5; 100.0%; Pred. No. 1.1e+05; tive 0; Mismatches 0; Indela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 AA; 626 MW; 6DD9C9CB10300000 CRC64;

        MOD_RES
        5
        AMIDATION.

        SEQUENCE
        5 AA; 656 MW; 71A9C9CB10300000 CRC64;

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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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                                                                                                                                                                                             SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE, AND MASS SPECTROMETRY.
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15-JUL-1998 (Rel. 36, Last annotation update) Triosephosphate isomerase (BC 5.3.1.1) (TIM)
                                                                      NCBI_TaxID=9615;
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                                                                                                                                                                                        phosphate,
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NON TER
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Suboesophageal ganglion pentapeptide.
Acheta domesticus (House cricket).
Bukaryota, Metazoa, Arthropoda, Mandibulata, Pancrustacea, Hexapoda,
Insecta, Perrygota, Neoptera, Orthopteroidea, Orthoptera, Ensifera,
Gryllidae, Gryllinae, Acheta.
                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                   Wabhitz P.A., Bowie J.H., Tyler M.J., Wallace J.C., "Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
                                                                                   Eukaryota; Metazoa; Chordata; Crāniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;0
                                                                                                                                                                                                      rubella.";
Aust. 'J. Chem. 52:639-645(1999).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
--- ACTIVITY.
--- TISSUE SPECIFICITY: SECRETED BY THE SKIN DORSAL GLANDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wicker C., Wicker C.;
"Isolation and structure of a peptide isolated from the suboesophageal ganglion of Acheta domesticus (orthoptera).";
Comp. Biochem. Physiol. 88C:185-187(1987).
-i- SUBCELLULAR LOCATION: MAIN PEPTIDE FROM THE SUBOESOPHAGEAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 26.1%; Score 6; DB 1; Length 5; Local Similarity 100.0%; Pred. No. 1.1e+05; es 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                            26.1%; Score 6; DB 1; Length 5; ilarity 100.0%; Pred. No. 1.1e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR; JS0319; JS0319. SEQUENCE 5 AA; 476 MW; 69D76DDDDDDDDD0000 CRC64;
                                                                                                                                                                                                                                                                        5 AA; 570 MW; 71A9C9C862A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                              01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
                             15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Litoria rubella (Desert tree frog)
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                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA.
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(Rel. 34, Last sequence update)
         5 AA.
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         STANDARD;
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                                                                                                                       NCBI_TaxID=104895;
                                                                                                                                                                                                                                                             Amphibian skin.
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ID TPIS'CANFA

AC P5471-1996 (

DT 01-0CT-1996 (
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P19991;
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Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACC clade;
Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Touzet P., Riccardi F., Morin C., Damerval C., Huet J.-C.,
Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                       Theor. Appl. Genet. 93:997-1005(1996).
-!- MISCELLANGOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 6.1, ITS MW IS: 30.4 kDa.
Maize-2DPAGE; P80628; COLEOPTILE.
                                                                                                                                                                                                                                                                                                                                                                                                                -!- PATHWAY: PLAYS AN IMPORTANT ROLE IN SEVERAL METABOLIC PATHWAYS. -!- SUBUNIT: HOMODIMER (BY SIMILARITY). -!- SIMILARITY: BELONGS TO THE TRIOSEPHOSPHATE ISOMERASE FAMILY. HSC-2DPAGE; P54714; DOG.
                                               Canis familiaris (Dog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000652; Triophos ismrse.
PROSITE; PS00171; TIM; PARTIAL.
Isomerase; Glycolysis; Gluconeogenesis; Fatty acid biosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .,
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01-OCT-1996 (Rel. 34, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Unknown protein from 2D-page of etiolated coleoptile (Spot 474)
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(TIM) (Fragment)
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MEDLINE=98163340; PubMed=9504812;
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                                                                                                                  Limulus polyphemus (Atlantic horseshoe crab), and Carcinus maenas (Common shore crab) (Green crab).

Eukaryota, Metazoa, Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blattoidea; Blattidae; Periplaneta.

NCBI_TaxID=6978, 6850, 6759;
                                                                                                                                                                                       SPECIES=P.americana;
MEDLINE=76074708; PubMed=576;
Starratt A.N., Brown B.E.;
"Structure of the pentapeptide proctolin, a proposed neurotransmitter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                      Stangier J., Dircksen H., Keller R.; "Identification of proctolin in
                                                                                                                                                                                                                                                              SPECIES=P.americana;
MEDILINESal122865; PubMed=6113690;
O'Shea M., Adams M.E.;
"Pencapeptide (proctolin) associated with an identified neuron.";
Science 213:567-569(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identification of proctolin in the central nervous system of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
0
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MEDLINE=90287800; PubMed=2356151;
Groome J.R., Tillinghast E.K., Townley M.A., Vetrovs A.,
Watgon W.H. III, Hunt D.F., Griffin P.R., Alexander J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                               21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
01-FEB-1995 (Rel. 31, Last annotation update)
Proctolin.
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                                                      5 AA.
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100.0%; Pred. No. 1.1
ive 0; Mismatches
                                                                                                         Periplaneta americana (American cockroach)
                                                                                                                                                                                                                                                                                                                                                                                  horseshoe crab, Limulus polyphemus.";
Peptides 11:205-211(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES=C.maenas;
MEDLINE=86232789; PubMed=2872661;
                                                                                                                                                                                                                                    Life Sci. 17:1253-1256(1975)
                                                      STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Conservative
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Matches 1; Conserv
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                                                     PRCT PERAM
P01373:
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PSK_DAUCA
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PATHWAY: Biotin biosynthesis.
SUBUNIT: HOMODIMER.
SIMILARITY: BELONGS TO CLASS-III OF PYRIDOXAL-PHOSPHATE-DEPENDENT
AMINOTRANSFERASES.
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Phytosulfokine-alpha (PSK-alpha) [Contains: Phytosulfokine-beta (PSK-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -!- SUBCELLULAR LOCATION: Secreted.
-!- PTM: SULFATION IS IMPORTANT FOR ACTIVITY AND FOR THE BINDING TO PUTATIVE MEMBRADE RECEPTOR (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE PHYTOSULFOKINE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=CV. US-Harumakigosun,
MEDLINE=20212743; PubMed=10750705;
Hanai H., Matsuno T., Yamamoto M., Matsubayashi Y., Kobayashi T.,
Kamada H., Sakagami Y.,
"A secreted peptide growth factor, phytosulfokine, acting as a
stimulatory factor of carrot somatic embryo formation.";
Plant Cell Physiol 41:27-32 (2000)
-!- FUNCTION: IN PRESENCE OF 2,4-D, STIMULATES PROLIFERATION OF THE
CELLS, BUT DOES NOT STIMULATE DIFFERENTIATION INTO THE SOMATIC
                                                                                                                                                                                              Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Asteridae, euasterids II, Apiales, Apiaceae, Daucus.
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01-JAN-1990 (Rel. 13, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Adenosylmethionine-8-amino-7-oxonomanoate aminotransferase
(RC 2.6.1.62) (7,8-diamino-pelargonic acid aminotransferase) (DAPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CATALYTIC ACTIVITY: S-adenosyl-L-methionine + 8-amino-7-oxononanoate = S-adenosyl-4-methylthio-2-oxobutanoate + 7,8-
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Pred. No. 1.1e+05;
2; Mismatches 0; Indels
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76C1BB504B300000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE, AND IDENTIFICATION BY MASS SPECTROMETRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PHYTOSULFOKINE-BETA
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MEDLINE=89006280; PubMed=2971595;
Shiuan D., Campbell A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COFACTOR: PYRIDOXAL PHOSPHATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17.48;
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les 0; Conservative
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                                                                                                                                                                      Daucus carota (Carrot).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diaminononanoate.
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                                                                                                                                                                                                                                                                                               NCBI_TaxID=4039;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDINE=89006280; PubMed=2971595;
Shiuan D., Campbell A.;
"Transcriptional regulation and gene arrangement of Escherichia coli,
"Citrobacter freundii and Salmonella typhimurium biotin operons.";
Gene 67:203-211(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- PATHWAY: Biotin biosynthesis; last step.
-!- SIMILARITY: BELONGS TO THE BIOTIN AND LIPOIC ACID SYNTHETASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JAN'1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
15-JUN-2002 (Rel. 14, Last annotation update)
Biotin synthase (EC 2.8.1.6) (Biotin synthetase) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 0.0%; Pred. No. 1.1e+65; nes 0; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                            Score 2; DB 1; Length 5;
Pred. No. 1.1e+05;
1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- CATALYTIC ACTIVITY: Dethiobiotin + sulfur = biotin.
                                                                                                                                                     EMBL; M21922; -; NOT_ANNOTATED_CDS.
InterPro; IPR000954; Aminotran 3.
PROSITE; PS00600; AA TRANSFER_CLASS 3; PARTIAL.
Biotin blosynthesis; Transferase; Aminotransferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 AA; 532 MW; 75A5B1EDD6F00000 CRC64;
                                                                                                                                                                                                                                                     NON TER 5 5 SEQUENCE 5 AA; 582 MW; GAAABIBIAGF00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; M21922; -; NOT_ANNOTATED_CDS.
Biotin biosynthesis; Tron-sulfur; Transferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                                                                                                                                                              8.7%;
0.0%; P
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Best Local Similarity 0.vv.,
0; Conservative
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                                                                                                                                                                                                                                       Pyridoxal phosphate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Citrobacter freundii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=546;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITFR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                    MEDLINE-88227859; bubMed=2836369; MEDLINE-88227859; bubMed=2836369; MEDLINE-88227859; bubMed=2836369; Inamoro S., Yoshioka Y., Ohtsubo B.; Indentification and characterization of the products from the traj and train genes of plasmid R100."; and train genes of plasmid R100."; J. Bacteriol. 170:2749-2757(1988).
I. FUNCTION: TRANSFER GENE PROTEIN. IS INVOLVED IN THE CONJUGATION PROCESS OF BACTERIAL GELLS FOR THE EXCHANGE OF PLASMID DNA.
I. SUBGELLULAR LOCATION: Cytoplasmic.
I. SIMILARITY: TO TRAM PROTEIN OF OTHER PLASMIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=L2/434/Bu;
Bini L., Santucci A., Magi B., Marzocchi B., Sanchez-Campillo M.,
Comanducci M., Christianen G., Birkelund S., Vtretou E., Ratti G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -i- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 4.5, ITS MW IS: 28 kDa.
Siena-2DPAGE; P38005; -.
NON TER
                                                                                                                                             Plasmid IncFII R100.
Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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01-0CT-1994 (Rel. 30, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Unknown protein from 2D-page from elementary body (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  y Match 0.0%; Score 0; DB 1; Length 5; Local Similarity 0.0%; Pred. No. 1.1e+05; hes 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chlamydia trachomatis.
Bacteria, Chlamydiales; Chlamydiaceae; Chlamydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SEP-1994) to the SWISS-PROT data bank
                                 01-JAN-1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; M20941; -; NOT_ANNOTATED_CDS.
PIR; A32014; A32014.
Conjugation; Plasmid; DNA-binding.
NON_TER
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STANDARD;
                                                                                          TraM protein (Fragment).
                                                                                                                               Escherichia coli.
                                                                                                                                                                                                      NCBI_TaxID=562;
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 TRM3 ECOLI
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Query Match

RESULT 19

Best Loďal Similarity 0.0%; Pred. No. 1.1e+05; Matches 0; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Search completed: February 12, 2003, 10:51:53 Job time : 12 secs

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Eukaryoča, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Archosauria, Aves, Neognathae, Galliformes, Phasianidae, Phasianinae,
                                                                                                                      PubMed=613771;
Dockray G.J., Reeve J.R. Jr., Shively J., Gayton R.J., Barnard C.S.;
"A novel active pentapeptide from chicken brain identified by
antibodies to PMRPamide.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hordeum vulgare (Barley).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, Pooldeae,
Triticeae, Hordeum.
NCBI_TaxID=4513;
                                                                                                                                                                                  Nature 305:328-330(1983).
-!- FUNCTION: MAY FUNCTION AS A NEUROTRANSMITTER OR MODULATOR.
-!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillales; Bacillaceae; Bacillus.
NCBL_TaxID=1396;
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                                                                                                                                                                                                                                                                                            h 26.1%; Score 6; DB 13; Length 5; Similarity 100.0%; Pred. No. 6.7e+05; 1; Conservative 0; Mismatches 0; Indels
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Last sequence update)
Last annotation update)
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01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA; 623 MW; 6B01AAA336F00000 CRC64;
                                                                                                                                                                                                                                                Neuropeptide.
SEQUENCE 5 AA; 645 MW; 69D4073767400000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Browne N., Dowds B.C.A., Submitted (JUL-2001) to the SWISS-PROT data bank. NON_TER \,5\,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   88 kDa protein (Fragment).
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Gallus gallus (Chicken).
                                                                                         SEQUENCE, AND SYNTHESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=NCIMB 11796;
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Best Local Similarity
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                                                        NCBI_TaxID=9031;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacillus cereus.
                                                                                                         TISSUE=BRAIN;
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                                                                                                   February 12, 2003, 10:48:54; Search time 28 Seconds (without alignments) 36.794 Million cell updates/sec
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              GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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01-UJN-2002 (TrEMBLrel. 21, Created)
01-UJN-2002 (TrEMBLrel. 21, Last sequence update)
01-UJN-2002 (TrEMBLrel. 21, Last annotation update)
FMRFamide-like neuropeptide (LPLRF-amide).
                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
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                                                                          OM protein - protein search, using sw model
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1: Sp_archea:*
2: sp_bacteria:*
3: sp_tungi:*
4: sp_tungi:*
5: pl_numan:*
5: sp_invertebrate:*
5: sp_mumal:*
5: sp_mumal:*
5: sp_mhc:*
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Gapop 10.0 , Gapext 0.5
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sp_virus:*
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Perfect score:
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Database :

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P83308

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RESULT 1 P83308

Result Š.

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Gaps

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0; Indels

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RC STRAIN=HIMALAYA; TISSUE=ALEURONE LAYER;
RA MEDLINE=9139704; Pubmed=1831055;
RT "Control of transient expression of chimaeric genes by gibberellic
RT acid and abscisic acid in protoplasts prepared from mature barley
RT aleurone layers."; 713-721(1991).
RL Plant Mol. Biol. 16713-721(1991).
CC -!- CATALYTIC ACTIVITY: ENDOHYDROLYSIS OF 1.4-ALPHA-GLUCOSIDIC
C. LINKAGES IN OLIGOSACCHARIDES AND POLYASACCHARIDES.
CC -!- MISCELLANEOUS: THERE ARE AT LEAST 4 TYPES OF ALPHA-AMYLASE IN
CC -!- MISCELLANEOUS: THERE ARE AT LEAST 4 TYPES OF ALPHA-AMYLASE IN
CC BARLEY.
CC --- MISCELLANEOUS: Carbohydrate metabolism; Seed; Germination;
KW Hydrolase; Glycosidase; Carbohydrate metabolism; Seed; Germination;
FT Calcium; Multigene family.
FT S SEQUENCE 5 AA; 600 MW; 61E3344DD6F00000 CRC64;
Ouery Match
Best Local Similarity 0.0%; Pred. No. 6 7e+05;
Matches 0; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
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Search completed: February 12, 2003, 10:52:27 Job time: 28 secs

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Endothelin C-termi
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Endothelin cyclic
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                 GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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Listing first 100 summaries
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SULT 1 733038 AAY3303	8 8 st.	T 1 038 AAY33038 standard; peptide;	pepti	de ;	ALIGNMENTS 5 AA.	

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This invention describes a novel method for preparing a peptide or recombinant antibody, which mimics an antigenic carbohydrate. The peptides and recombinant antibodies prepared to mimic antigenic carbohydrates can be used to enhance binding of anti-antigenic carbohydrate antibodies to the antigenic carbohydrate, e.g. as vaccine carbohydrate antibodies to the antigenic carbohydrate, e.g. as vaccine adjuvants. The peptides to be used to inhibit binding of a ligand to a receptor, which is an antigenic carbohydrate. The methods are used to prepare the peptides and antigenic antibodies, which mimic the antigenic carbohydrates. Carbohydrate-conjugate vaccines could be used in tumour immunity and cancer therapy. The peptides and antibodies can also be used as antibacterials. Peptides that mimic carbohydrate antigens can be carbohydrate antigens can be committed to develop a longer lasting immune response. Other advantages of the peptides can be precisely defined; (2) the immunogenicity of the peptides can be precisely defined; (2) the immunogenicity of the peptides can be significantly enhanced by polymerization or addition or relatively small carrier molecules that reduce the total
                                                                                                                                                                                                         Mimotope, antigen, carbohydrate, antibody, mimic; vaccine, adjuvant, tumour immunity, cancer therapy, antibacterial, immune response, immunogenicity, anti-idiotype, T cell response manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptides and recombinant antibody mimics of carbohydrate antigens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       used for, e.g. treatment of cancer and infection
                                                                                                                                                                 Carbohydrate antigen peptide mimotope 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 21; Page 72; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US02405.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0073690.
                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kieber-Emmons T;
                                                                                                                                                                                                                                                                                                                                     WO9940433-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-1999;
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                                                                                                                         03-NOV-1999
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                                                                                  AAY33038;
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Tumour metastasis requires detachment of malignant cells from the primary tumour, penetration of blood or lymph vessels and attachment to the endothelium of distant organs, ultimately resulting in the formation of new tumours. The selectin family of adhesion molecules is implicated in this process. E-selectin family of adhesion molecule significated in this process. E-selectin family of adhesion molecule spreased by activated vascular endothelium. E-selectins bind to glycoconjugates carrying a terminal tetrasaccharide Lewis antigen, which are found on tumour cell surfaces. One such Lewis antigen is Lewis Y (LeY). The binding of selectin molecules to their ligands is thought to be an important step in metastasis. Therefore, inhibition of E-selectin-dependent cancer therapy. The present sequence is a non planar-X-planar type motif. This motif is thought to bind to anti-LeY antibody BRS5-2. BRS5-2 is a peptido-mimetics of E-selectins. Peptides that block E-selectin-LeY binding and therefore adhesion of tumour cells and leukocytes to endothelial cells inhibit metastasis.
                                                                                                                                                                                                                               0
amount of antigen required for immunization; (3) peptide synthesis may be more practical than synthesis of carbohydrate-protein conjugates or the production of anti-idiotypes; (4) peptide minicking sequences can be engineered into DNA plasmids for DNA vaccination to further manipulate T cell responses. ANY33028-Y33308/ represent the carbohydrate antigen peptide mimotopes described in the invention.
                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                       87.0%; Score 20; DB 20; Length 5; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Motif binding to anti-Lewis antigen anibody BR55-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (WIST-) WISTAR INST ANATOMY & BIOLOGY. (UYPE-) UNIV PENNSYLVANIA.
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                                                                                                                                                                                                                                                                                                                                                                                                             AAB07280 standard; peptide; 5 AA.
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                                                                                                                                                                                                          Local Similarity 60.0 ies 3; Conservative
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                                                                                                                                                  5 AA;
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DB 21; Length 5;

87.0%; Score 20;

5 AA;

Sequence Query Match

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                       Gaps
                                                                                                                                                                                                                                                                                                                                                                      precocious puberty; amenorrhoea; premenstrual syndrome; cyclic; multilocular ovarian syndrome; comedo; pregnancy; contraception; ovulation; menstrual cycle; oestrus; meat quality; growth; spawning; fish; testosterone; superagonist; leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "N-pToluenesulphonylarginine, D-form residue"
                                                                                                                                                                                                                                                                                                               Luteinising hormone releasing hormone receptor; LH-RH; antagonist;
sex-hormone; human; veterinary medicine; cancer; prostate; uterus;
breast; pituitary; prostatomegaly; endometriosis; hysteromyoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclic penta:peptide(s), some new, as LH-RH receptor antagonists used to treat or prevent sex hormone dependent disorders, e.g. cancer, also for control of pregnancy and menstruation and to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'note= "Joined via a peptide linkage to Phel"
                                                                                                                                                                                                                                                                              Cyclic pentapeptide #24 used in LH-RH receptor antagonist.
Pred. No. 7.8e+05;
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                     0; Mismatches
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                                                                                                                                                                         AAW17808 standard; peptide; 5 AA.
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 60.0%;
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                    3; Conservative
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Best Local Similarity
Matches 3; Conser
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                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
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100.0%; Pred. No. 7.8e+05;
vative 0; Mismatches 0;
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                                                                                                                                                                                   AAR89911 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                 p53/MDM2 binding inhibitor #1.
                                     82.68;
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94US-0277660.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95WO-GB01719
                                                                                                                                                                                                                                       10-SEP-1996 (first entry)
                                     Query Match 82.6
Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYDU-) UNIV DUNDEE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference 3
             5 AA;
                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9602642-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-FEB-1996
                                                                                          1 FXXLW
                                                                                                                   1 FRFLW
                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                              AAR89911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                       AAR89911
X S
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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                p53 tumour suppressor procein and the MMM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptides of the invention which interfere with the binding of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     82.6%; Score 19; DB 17; Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 19; _____Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human p53 consensus hydrophobic interaction motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                        /note= Any amino acid
                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY58280 standard; peptide; 5 AA.
                           AAR89912 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 29; 46pp; English.
                                                                                                                   p53/MDM2 binding inhibitor #2.
                                                                                                                                                                                                                                                                                                                                                95WO-GB01719.
                                                                                                                                                                                                                                                                                                                                                                             95US-0424957.
                                                                                                                                                                                                                                                                                                                                                                                        94US-0277660.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     80.08;
                                                                                      10-SEP-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAR-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                        (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           other malignancies
                                                                                                                                                                                                                                         Misc-difference 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                           20-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                              20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                             19-APR-1995;
                                                                                                                                                                                                                                                                                     WO9602642-A1
                                                                                                                                                                                                                                                                                                                  01-FEB-1996
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                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence
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                                                         AAR89912;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lane DP,
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RESULT 5
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The invention relates a method of identification of a compound that

modulates coactivator binding to a nuclear receptor. The method

comprises modelling test compounds that fit spatially into a nuclear

receptor (NR) coactivator binding site of interest using an atomic

structural model of a nuclear receptor coactivator binding site. The

coactivator binding site, leading to the interest using an atomic

coactivator binding site, leading to the indentification of a compound

which modulates coactivator binding. Nuclear receptors are a superfamily

coactivator binding site, retinoid receptors (RARs and RARs), peroxisome

commone receptors (TRR), retinoid receptors (RARs and includes thyroid

copperors (PRARs), quecocrticoid receptors (RRS), peroxisome

receptors (PRARs), quecocrticoid receptors (RRS), progestin receptors

copperors (RRS), These receptors (RRS), androgen receptors

construction, for example NRS can stimulate transcription in

compare ligand, also bind coactivators proteins that are involved in

compare ligand, also bind coactivators of the plob (Emily mediate

compare to hormone binding by recruiting coactivator proteins to the

compare receptor's ligand binding domain, AP2, that is part of the

conclusive of a transcriptional activation domain, AP2, that is part of the methods are appoint or antelear receptor's ligand binding domain. The method may be used to

compare the methods can also be used to identify residues which

men methods are an also be used to identify residues which

men methods are an also be a nuclear receptor of interest.
Nuclear receptor; pl60 coactivator; NR-box; coactivator binding domain; NR; AF2 transactivation domain; wild-type; thyroid hormone receptor; TR; retinoid receptor; RAR; RXR; peroxisome receptor; PPAR; RXR; peroxisome receptor; PR; vitamin D receptor; PR; ostrogen receptor; PR; mineralocorticoid receptor; MR; androgen receptor; AR; Grip-1; Tif2; NcoA-2; RAC3; AIB1; TRAM-1; P/CIP; SRC1; breast cancer; prostate cancer; cardiac arrhythmia; infertility; osteoporosis; hyperthyroidism; hypercholesterolaemia; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The methods are applicable to generating new compounds that distinguish nuclear receptor isoforms. This can facilitate generation of either rissue specific or function-specific compounds. Nuclear receptors have been implicated in a variety of medical disorders, including breast cancer, prostate cancer cardiac arrhythmia, infertility, osteoporosis, hyperthyroidism, hypertholesterolomema and obesity. Sequences AAV58282-AAV58302 represent the regions (NR-boxes 1-3) of p160 coactivator proteins which interact with the nuclear receptor coactivator binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying modulators of nuclear receptor coactivator binding useful for generating new compounds which distinguish nuclear receptor isoforms -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fletterick R, Kushner PJ;
                                                                                                                                                                                                                                                                                                                                                          'note= "Xaa = any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Xaa = any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 14B; Page 12; 281pp; English.
                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Darimont B, Feng W, F
West BL, Yamamoto KR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US06899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0113146.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         98US-0079956.
                                                                                                                                                                                                                                                                                                                                         'label= Xaa
                                                                                                                                                                                                                                                                                                                                                                                                            /label= Xaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-072429/06.
                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference 3
                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAR-1999;
                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9960014-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-NOV-1999.
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Wagner RL,
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sites (AAYS8307-Y58328). The NR coactivator binding sites comprise two regions of the NR; helices 3-6 (H3-H6), and helix 12 (H12). The coactivator NR-boxes all share a consensus sequence LXXLL (AAYS8278), with the NR-box 2 motifs sharing the consensus motif ILXXLL (AAYS8279). Sequences AAYS8280-Y58281 represent the hydrophobic interaction motifs of p53 and VP16 used in an exemplification of the invention. Sequences AAYS8329-Y58240 represent mutant versions of the human Grip-1 NR-box 2 sequence (AAYS8283), and sequences AAYS8341-Y58353 represent mutant versions of NR coactivator binding sites, used in exemplifications of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cytostatic; cardiovascular; antiarrhythmic; vasotropic; osteopathic; endocrinal; antilipaemic; anorectic; coactivator binding; lead compound; nuclear receptor; NS; peptidomimetic; scaffold; breast cancer; nuclear receptor-based disorder; prostate cancer; cardiac arrhythmia; infertility; osteoporosis; hyperthyroidism; hypercholesterolaemia; obesity; ExxLW motif; p53; glucocorticoid; GR; protein co-ordinate data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kushner PJ;
                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Method for identifying agonist and antagonist compounds used for mobilating nuclear receptor coactivator binding, which are useful treating nuclear receptor-based disorders e.g. breast cancer, infertility, or hyperthyroidism -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arnold JR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Feng W, Fletterick RJ,
, Geistlinger TR, Arnol
                                                                                                                                                                                                      82.6%; Score 19; DB 21; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                            100.0%; Prec. ....
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hydrophobic interaction FxxLW motif of p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "Xaa is unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /label= unknown
/note= "Xaa is unknown"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 14; Page 15; 307pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Darimont B, F
Yamamoto KR,
                                                                                                                                                                                                                                                                                                                                                                      AAO20822 standard; Peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 label= unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JUN-2001; 2001WO-US20969.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JUN-2000; 2000US-0609361
                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Baxter JD,
,, West BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-315164/35.
                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                       5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
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                                                                                                                                                                                                                                                               1 FXXLW 5
                                                                                                                                                                                                                                                                                           1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-JAN-2002
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                                                                                                                                           invention.
                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                     AA020822;
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                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Guy RK,
                                                                                                                                                                                                                               Matches
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The invention relates to a method for identifying a compound that

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modulates coactivator binding to a nuclear receptor (NR). The method comprises modelling test compounds that fit electrostatically into an NR caactivator binding site (BS) using an appropriate atomic structural model of (BS); screening the test compounds in an assay characterised by binding of a test compound to (BS); and identifying a library or test compound that modulates coactivator binding to the NR. The methods of the compound that modulates coactivator binding to the NR. The methods of the invention are useful for identifying peptides, peptidomimetics, or small cativity, particularly new lead compounds, scaffolds and combinatorial libraries. The methods are used in selecting and characterising structure (function relationships of natural or synthetic coactivator compounds, which are useful in treating nuclear receptor-based disorders e.g. breast cancer, prostate cancer, cardiac arrhythmia, infertility, osteoporosis, hyperthyroidism, hypercholesterolaemia or obesity. This sequence corpusation (RR) nuclear receptor of the invention is able to interact context in motific with peptides containing this motif.
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "(Ac- or 1-adamantyl acetic acid)- D-Phe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hypertension, myocardial infarction; congestive heart failure; endotoxic shock; subarachnoid haemorrhage; asthma; arrhythmias; acute renal failure; preeclampsia; diabetes; metabolic; endocrinological; neurological; disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide(s) used as endothelin antagonists - for treating hypertension, metabolic and endocrine disorders, heart failure,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The peptide is an endothelin antagonist useful in controlling hypertension, myocardial infarction, congestive heart failure,
                                                                                                                                                                                                                                                                                                                                                                        82.6%; Score 19; DB 23; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diabetes, asthma, neurological disorders, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Depue P, Doherty AM, Taylor MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR29438 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Endothelin antagonist peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92WO-US03408,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91US-0701274.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91US-0809746
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-APR-1993 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                5; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1992-415706/50.
                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR29438;
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                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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             acute renal failure, precclampsia, diabetes and metabolic, endocrinological and neurological and neurological disorders. Administration is oxal parenteral or by inhalation in doses of 0.01-20 (esp. 0.01-10) mg/kg/day. It may be prepared by conventional peptide synthesis.
                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide(s) used as endothelin antagonists - for treating hypertension, metabolic and endocrine disorders, heart failure, diabetes, asthma, neurological disorders, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                           Hypertension; myocardial infarction; congestive heart failure; endotoxic shock; subarachnoid haemorrhage; asthma; arrhythmias; acute renal failure; preeclampsia; diabetes; metabolic; endocrinological; neurological; disorders.
endotoxic shock, subarachnoid haemorrhage, asthma, arrhythmias,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The peptide is an endothelin antagonist useful in controlling
                                                                                                                             Length 5;
                                                                                                                                                           2; Indels
                                                                                                                           Query Match 73.9%; Score 17; DB 13; Best Local Similarity 40.0%; Pred. No. 7.8e+05; Matches 2; Conservative 1; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Taylor MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Ac-D-Phe"
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                                                                                                                                                                                                                                                                                                      AAR29448 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Doherty AM,
                                                                                                                                                                                                                                                                                                                                                                                                   Endothelin antagonist peptide.
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Best Local Similarity 40.0%
Best Local 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                5 AA;
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                                                                                                                                                                                             1 FXXLW 5
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1 FDIIW 5
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                                                                                                Sequence
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endocrine disorders, congestive heart failure, myocardial infarction, endotoxic shock, subarachnoid haemorrhage, arrhythmia, asthma, acute and chronic renal failure, preeclampsia, diabetes, neurological disorders, pulmonary hypertension, ischaemic disease, ischaemic bowel disease, gastric mucosal damage, Raynaud's disease, restenosis, percutaneous transluminal coronary angioplasty, angina and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel antagonists of endothelin are claimed which are C-terminal hexapeptides and analogs of BT-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example of the generic compounds.

The peptides are useful for treating hypertension, metabolic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina, cancer etc.
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                                                                                                                                                                                         Endothelin C-terminal peptide analog, useful as antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73.9%; Score 17; DB 15; Length 5; 40.0%; Pred. No. 7.8e+05; tive 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         He JX, Taylor MD;
                                                                                                                                                                                                                       Endothelin; ET-1; receptor; antagonist.
                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 5; Page 119; 146pp; English.
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                                                                                                                                                                                                                                                                                                            /note= "Ac-D-Phe"
                                                                                                   AAR69220 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cody WL, Depue P, Doherty AM,
                                                                                                                                                                                                                                                                                                                                                                                                  93WO-US12377.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (WARN ) WARNER LAMBERT CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1994-234617/28.
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les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
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1 FLDIW 5
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1 FXXLW 5
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                         1 FLDIW
                                                                                                                                                                                                                                                   Synthetic.
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                                                                     RESULT 10
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Score 17; DB 13; Length 5; Pred. No. 7.8e+05; 1; Mismatches 2; Indels

73.9%;

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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= D-Glu
                                                               93WO-US12377
                                                                                   92US-0995480
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                                                                                                     (WARN ) WARNER LAMBERT
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les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                5 AA;
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                                                               17-DEC-1993;
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                         WO9414843-A.
                                            07-JUL-1994.
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cancer etc.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                            Novel antagonists of endothelin are claimed which are C-terminal hexapeptides and analogs of BT-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example
                                                                                                                                                                                                                                                               New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina,
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                   Endothelin C-terminal peptide analog, useful as antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin C-terminal peptide analog, useful as antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  73.9%; Score 17; DB 15; Length 5; 40.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                           Taylor MD;
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                                                                                                                                                                                                                           не лх,
                                        Endothelin; ET-1; receptor; antagonist
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                                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                                                                        Claim 5; Page 120; 146pp; English.
                                                                                                    /note= "Ac-D-Phe"
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                                                                                                                                                                93WO-US12377
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07-MAR-1995 (first entry)
                                                                                                                                                                                                     (WARN ) WARNER LAMBERT CO
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                                                                                                                                                                                                                                                                                                                                                                     of the generic compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Conservative
                                                                                                                                                                                                                                             WPI; 1994-234617/28
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Matches 2; Conserv
                                                                                         Misc-difference 1
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Misc-difference 1
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1 FDFIW 5
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                                                                                                                       WO9414843-A
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                                                                                                                                                                                                                                                                                      cancer etc.
                                                            Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                         Cody WL,
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 133X8X8X8X8X1
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of the generic compounds.

The peptides are useful for treating hypertension, metabolic and endocrine disorders, congestive heart failure, myocardial infarction, endotoxic shock, subarachnoid haemorrhage, arrhythmia, asthma, acute and chronic renal failure, preeclampsia, diabetes, neurological disorders, pulmonary hypertension, ischemic disease, ischaemic bowel disease, gastric mucosal damage, Raynaud's disease, restenosis, percutaneous transluminal coronary angioplasty, angina and cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel antagonists of endothelin are claimed which are C-terminal hexapetides and analogs of ET-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Taylor MD;
1-adamantyl-acetyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      не лх,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /label= D-aile, D-val
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Gaps

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The peptide is prepd. by cyclisation of the synthetic peptide. The Trp residue at posn. 3 may be side chain protected with a lower alkanoyl gp; the Glu residue at posn. 4 may be protected with an al (lower) alkyl gp. An analogous natural peptide, WS 7338, may be isolated from cultures of Streptomyces sp. no. 7338.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide prepd. by culturing Streptomyces spp - used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 5;
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                                                             Length 5;
                                                                                  0; Indels
isolated from cultures of Streptomyces sp. no. 7338.
See also AAR13339 and AAR13909.
                                                           65.2%; Score 15; DB 12; L
100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
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                                                                                                                                                                                         AAR13909 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                             Endothelin antagonist peptide
                                                                                                                                                                                                                                                                                                                                                        label= D-Val
                                                                                                                                                                                                                                                                                                                                                                               /label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                     /label= D-Glu
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                                                                      Local Similarity 100.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               endothelin antagonist
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les 2; Conserva
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                                       5 AA;
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                                        Sequence
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                                                             Query Match
                                                                                                                                                                                                                                                                                     Cyclic.
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AAR38437
                                                                                     Matches
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                                                                                                                                                     The Trp residue at posn. 3 may be side chain protected with a lower alkanoyl gp; the Glu residue at posn. 4 may be protected with an al (lower) alkyl gp. An analogous natural peptide, WS 7338, may be isolated from cultures of Streptomyces sp. no. 7338.
                                                                                                                                               peptide is prepd. by cyclisation of the synthetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide prepd. by culturing Streptomyces spp - used as endothelin antagonist.
                                                                                     New peptide prepd. by culturing Streptomyces spp - used as endothelin antagonist.
                                                                                                                                                                                                                                                 65.2%; Score 15; DB 12; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
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                                                                                                                         Claim 1; page 1; 15pp; Japanese.
                                                                                                                                                                                                      See also AAR13909 and AAR14025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'label= D-alle
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endothelin antagonist peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label= D-Trp
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                  89GB-0019726.
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                                        (FUJI ) FUJISAWA PHARM KK.
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Best Local Similarity
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                                                               WPI; 1991-241129/33.
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                  31-AUG-1989;
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The sequences given in AAR38436-38 are cyclic peptides which act as endothelin receptor antagonists. They may be used in the treatment of cardiac disease, acute renal insufficiency or asthma.
                                                                                                                                                                                                                                                        Novel cyclic peptide(s) used as endothelin receptor antagonists for treating cardiac diseases, e.g. myocardial infarction, acute renal insufficiency or asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /label= OTHER
/note= "alpha-Abu, delta-Ava or epsilon-Aca"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cyclic, endothelin receptor; antagonist, cardiac disease, myocardial infarction; acute renal insufficiency; asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin receptor antagonist peptide #1.
               'note= "D-form residue"
                                                                                             /note= "D0form residue"
                                                                      'note= "D-form residue"
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                                                                                                                                                                                                                                                                                                        Disclosure; Page 6; 8pp; Japanese.
                                     /label= OTHER
/note= "beta-Asp"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR38436 standard; peptide; 5 AA.
                                                                                                                                                                                      91JP-0191454.
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Best Local Similarity 100.0
Matches 2; Conservative
                                                                                                                                                                                                                                    WPI; 1993-282913/36.
   Misc-difference 1
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                                                                                                                                                                                                                                                                                                                                                                                                                  Novel cyclic peptide(s) used as endothelin receptor antagonists - for treating cardiac diseases, e.g. myocardial infarction, acute renal insufficiency or asthma
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/note= "alpha-Abu, delta-Ava or epsilon-Aca"
                                                                                        Cyclic, endothelin receptor, antagonist, cardiac disease, myocardial infarction, acute renal insufficiency, asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 14; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
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                                                                  Endothelin receptor antagonist peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Endothelin receptor antagonist peptide #3.
                                                                                                                                                                                                                                'note= "D-form residue"
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 AAR38437 standard; peptide; 5 AA
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                                                                                                                                                                                    Misc-difference 2
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                      AAR38437;
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action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase
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                                                                                                                                                                                                                AAR35419 standard; peptide; 5 AA.
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                                                                                                                2; Conservative
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                                                                                                    Best Local Similarity
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                                              synthesis.
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                                                                                                                                                          The sequences given in AAR38436-38 are cyclic peptides which act as endothelin receptor antagonists. They may be used in the treatment of cardiac disease, acute renal insufficiency or asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
                                                                                           Novel cyclic peptide(s) used as endothelin receptor antagonists -
for treating cardiac diseases, e.g. myocardial infarction, acute
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                                                                                                                                                                                                                             Length 5;
                                                                                                                                                                                                                            65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Endothelin receptor inhibitor peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'note= "D-form residue"
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                                                                                                                                       Example 1-3; Page 6; 8pp; Japanese.
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                                                                                                                   renal insufficiency or asthma
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                           91JP-0191454.
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     92JP-0173065.
                                                (TAKE ) TAKEDA CHEM IND LTD.
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Best Local Similarity 100.v
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                                                                                                                                                                                                       5 AA;
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     30-JUN-1992;
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65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
       Length 5;
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Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 2; Conservative 0; Mismatches 0;
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JP05059098-A.
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                                                            Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  The sequences given in AAR35418-28 are cyclic peptides which act as endothelin inhibitors. They strongly inhibit the vasoconstrictive action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase
                                                                                                                                                                                                                                                                                                                                                                     New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                                                                                                                   Endothelin receptor inhibitor peptide #4.
                                                                                                                                                                                                     /note= "D-form residue"
                                                                                                                                                                                                                                          /note= "D-form residue"
                                                                                                                                                                                                                        'note= "D-form residue'
                                                                                                                                                                                    Location/Qualifiers
                                                             AAR35421 standard; peptide; 5 AA
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Matches 2; Conserv
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                                          RESULT 2
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                                             Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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100.0%; Pred. No. 7.8e+05;
tive 0; Mismatches 0;
Endothelin receptor inhibitor peptide #5.
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                                                                                                                                                                                                  'note= "D-form residue"
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Misc-difference 3
                                                                                                                                                       Location/Qualifiers
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91JP-0220610.
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(TAKE ) TAKEDA CHEM IND LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                                New cyclic penta:peptide derivs. contg. one L-aminoacid --inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                                                                                                                                                                                                                                                                                                      Length 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endothelin receptor inhibitor peptide #7.
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                      /note= "D-form residue"
    /note= "D-form residue"
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nes 2; Conserva
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New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endothelin receptor inhibitor peptide #8.
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                                                                RESULT 27
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endothelin inhibitors. They strongly inhibit the vasoconstrictive action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase synthesis.
                                                                                                                                                                                                                                                                                              Cyclic, endothelin, inhibitor; vasoconstriction, drug, circulation, mammal; myocardial infarction; renal dysfunction, asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endochalin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and asthma, etc.
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                                                                                           65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; rative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                           Endothelin receptor inhibitor peptide #9.
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                                                                                                                 Conservative
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                                                                                                                                                                                       Cyclic, endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
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Pred. No. 7.8e+05;
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7 100.0%; Pred. No. /...
7 0; Mismatches
                                                                                                                                          Endothelin receptor inhibitor peptide #10.
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AAR35427 standard; peptide; 5 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin; metal conjugate; complex; diagnostic; contrast agent;
X-ray imaging; NMR imaging; pathological vascular change; stenolic;
atherosclerosis; radiopharmaceutical; non-invasive diagnosis;
arterial wall lesion; pharmacokinetic.
                                      Cyclic, endothelin, inhibitor, vasoconstriction, drug, circulation, mammal; myocardial infarction; renal dysfunction, asthma.
                                                                                                                                                                                                                                                                                     New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                  Endothelin receptor inhibitor peptide #11.
                                                                                                         'note= "D-form residue"
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                                                                                                                             note= "D-form residue"
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                                                                                      Location/Qualifiers
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Matches 2; Conserv
                                                                                               Misc-difference 1
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                                                                                                                   Misc-difference 3
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This invention describes novel complexes of compounds which have the formula B-L-(K)b with metal ions of atomic number 21-32, 37-39, 42-51 and 57-38 where E = residue of an endothelin (or its derivative, partial sequence, analogue or antagonist), including forms with free SH groups that bind directly to metal, optionally labelled with radioactive iodine isotopes, L = bond, Z1-R-Z2 or a group (i), R = 1-20. Optionally constituted by 10 and/or epoxy, and coptionally interrupted by 1 or more 0, S, CO, NCO or NH (the last 2 optionally interrupted by 1 or more 0, S, CO, NCO or NH (the last 2 optionally nubstituted by 1-6C alkyl), Z1 and Z = 0, S, COO, NH, CS, NH, NGO or NHCS, s and t = 0-3, B = phenyl or cyclohexyl ring, b = 0 or 1, K is a chetating residue. The compounds of the invention are useful as diagnostic agents, including contrast agents for X-ray and NMR imaging of pathological vascular changes e.g.

CX ray and NMR imaging of pathological vascular changes before these become standiopharmaceuticals. The invention allows the marking of possible early non-invasive diagnostic agents they have better pharmacokinetic properties and provide improved contrast. They have better pharmacokinetic properties and provide improved contrast. They have better pharmacokinetic properties and provide improved contrast. They have excellent in vitro and in vivo stability on significant release or exchange of bound radionucleotides). This sequence is a specific claimed example of a peptide used in the construction of the metalcomplex-conjugates of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New complexes of derivatised endothelin or related cpds. - with radioactive or paramagnetic metal ions, useful as diagnostic agents for imaging pathological changes in blood vessels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kramp W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hilger CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                               'note= "Residue is alloDIle"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , Gries H, Hilger Ct
Speck U, Reiser JH;
                                                                                                                                      /note= "D-form residue"
/note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DIAG-) INST DIAGNOSTIKFORSCHUNG GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Page 31; 40pp; German.
                                                                                                                                                                                                                                                                                                                                                    93DE-4301871.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Erber S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schier H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
nes 2; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DINKELBORG L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1994-226321/28.
                                                                                                     Misc-difference 5
                             Misc-difference 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Dinkelborg L,
                                                                                                                                                                                                                                                                                                                                                    13-JAN-1993;
                                                                                                                                                                                                              DE4301871-A1
                                                                                                                                                                                                                                                                                                                                                                                                                        13-JAN-1993;
                                                                                                                                                                                                                                                                                 14-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Platzek J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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immunosuppression; CD4; gp120; T cell activation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                          The peptide inhibits binding of endothelin-1 (ET) to ETA receptors and/or to ETB receptors. It can be used for detecting, distinguishing and isolating ET receptor subtypes. The peptides also act as ET antagonists and can be used to treat endothelinmediated disorders such as hypertension, pulmonary hypertension, cardiovascular disease, bronchoconstriction, asthma, inflammatory diseases, ophthalmolgic shock, anaphylactic shock, haemorrhagic shock, gastroenteric disease, renal failure, endotoxin shock, menstrual disorders, obstretic conditions, erythropoietin-mediated vasoconstriction and wounds.
                                                                                                                                                                                                                                                                                                                                                              New cyclic peptide(s) which modulate endothelin activity - used for treating endothelin-mediated disorders such as cardiovascular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monomeric peptide #1, autoimmunogenic in HIV-infected subjects.
                               isolation; antagonist; hypertension; pulmonary hypertension; cardiovascular disease; bronchoconstriction; asthma; inflammatory disease; bronchoconstriction; anaphylactic shock; haemorrhagic shock; gastroenteric disease; renal failure; endotoxin shock; menstrual disorders; obstretic conditions; erythropoietin-mediated vasoconstriction; wound treatment.
                                                                                                                                                                           /note= "D-Trp bonded to D-Asp at position 1."
                      Cyclic; peptide; endothelin; inhibitor; receptor; detection;
                                                                                                                               'note= "D-Asp bonded to D-Trp at position
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1; HIV-2; human immunodeficiency virus; AIDS; acquired immune deficiency syndrome; autoimmune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
Cyclic peptide which modulates endothelin activity.
                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR46691 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 51; 58pp; English.
                                                                                                                                                                                                                                                                                               (IMMU-) IMMUNOPHARMACEUTICS INC
                                                                                                                                                     /label= D-His.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                     and respiratory diseases
                                                                                                                                                                                                                                                                                                                     Chan MF;
                                                                                                                                                                                                                                                                                                                                          WPI; 1994-007458/01.
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                                                                                                                   Misc-difference 1
                                                                                                                                          Misc-difference 3
                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 AA;
                                                                                                                                                                                                                                         17-JUN-1993;
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                                                                                                                                                                                                                    23-DEC-1993,
                                                                                                                                                                                               W09325580-A
                                                                                                                                                                                                                                                                                                                     Balaji VN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LW 5
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New peptide(s) inducing an auto immune response in HIV infected subjects - also the multimer, proteins contg. them and derived antibodies, for treating or preventing HIV infection and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
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                                                                                                                                                                                                                                                                                                 /note= "C-terminus is opt. extended
by 1-5 amino acids"
                                                                                                                                                                                                         /note= "N-terminus is opt. extended
by 1-5 amino acids"
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                                                                              Human Immunodeficiency Virus type 1
                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR60079 standard; Protein; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93WO-FR00803
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93FR-0001628
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zagury J, Zagury JF;
therapy; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1994-065602/08.
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(ZAGU/) ZAGURY J F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                              Misc-difference 5
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                                                                                                                                                                         Misc-difference
                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .0-AUG-1992;
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The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cyclic pentapeptides, useful as endothelin receptor antagonists are given
                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                              The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cycl penteapeptides, useful as endothelin receptor antagonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endothelin; ET; ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                                                                                                                                                                                                                                                                                                                Ouery Match
65.2%; Score 15; DB 15; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                                                                Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IAN-1477D - cyclic ET receptor antagonist.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                 Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR60082 standard; Protein; 5 AA.
91JP-0148807.
                                91JP-0148807.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            91JP-0148807
                                                                (TAKE ) TAKEDA CHEM IND LID.
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                                                                                                 WPI; 1994-260512/32
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                                                                                                                                                                                                                                                                                                                     5 AA;
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20-JUN-1991;
                                20-JUN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR60082;
                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cycl pentapeptides, useful as endothelin receptor antagonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 15; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Endothelin; ET; ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                       Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TAN-1462B - cyclic ET receptor antagonist.
                                                                                                 /label= alle
/note= "D-form residue"
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                                                                                                                                                   /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note= "D-form residue"
                                                                   'note= "D-form residue"
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                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR60080 standard; Protein; 5 AA.
                                                                                                                                                                                                                                                     91JP-0148807.
                                                                                                                                                                                                                                                                                       91JP-0148807.
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Rest Local Similarity 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                   Misc-difference
                                                                                                                                     Misc-difference
                                                                                   Modified-site
                                                                                                                                                                                                                                                                                       20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP06192293-A.
                                                                                                                                                                                                                                                     20-JUN-1991;
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                                                                                                                                                                                                                       12-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 LW 5
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   Synthetic
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          RESULT 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cyclic
                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pentapeptides, useful as endothelin receptor antagonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                       Endothelin, ET; ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                   65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              65.2%; Score 15; DB 15; L
100.0%; Pred. No. 7.8e+05;
ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                       /note= "D-5-methyltryptophan"
                                                                                                                                                                                                     TAN-1477K - cyclic ET receptor antagonist.
                                                                                                                                                                                                                                                                                                                      'note= "D-form residue"
                                                                                                                                                                                                                                                                                           'note= "D-form residue"
                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                  AAR60088 standard; Protein; 5 AA.
                                                                                                                                                                                                                                                                                                             /label= alle
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                                                                                                                                                                                     (first entry)
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-260512/32.
                                                          Best Local Similarity
Matches 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                   5 AA;
                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AA;
                                                                                                                                                                                                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                     21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                         20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                            20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                       JP06192293-A
                                                                                                                                                                                                                      Endothelin;
                                                                                                                                                                                                                                                                                                                                                                         12-JUL-1994.
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                                                                                     4 LW 5
                                                                                                     4 LW 5
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                                   Sequence
                                                                                                                                                                   AAR60088;
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                                                                                                                                RESULT 35
                                                                                                                                         AAR60088
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                                                                                                                                                                                                 CENP-B protein; centromere protein; epitope; autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; SRV-1 viral protease; systemic lupus erythematosus; scleroderma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A comparison of the CENP-B centromere protein sequence with proteins from immunoinfective cluster viruses revealed widespread homologies. The importance of these homologies motifs is that they are epitopes for autoantibodies occurring in high titres in systemic rheumatic disorders. Sera from such patients could be used for treatment of immunoinfective cluster virus infections, e.g. HIV-1, immunoinfective adenoviruses, human lymphotropic retroviruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                CENP-B protein residues 191-195, homologous to SRV-1 protease motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Methods for treating immunoinfective cluster virus infections -
utilise antibodies or fragments characteristic of auto antibodies
produced by patients with rheumatic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rubella virus, CMV and EBV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Takehana Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 71; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR71907 standard; peptide; 5 AA.
AAR62195 standard; Protein; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYSC-) UNIV SOUTHERN CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94WO-US02631.
                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Douvas A, Ehresmann G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1994-302689/37.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                 03-MAY-1995
                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                           WO9420141-A.
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                                                AAR62195;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR71907;
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with the C-terminal amino acid"
                                                                                                                                                                                                                                                                                                                                                                                     Wakimasu M;
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                                    /note= "Ser(Bzl)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 10; Page 16; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR69267 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                   Kikuchi T,
                                                                           /note= "D-Leu"
                                                                                                                                                                                                                                                                                                           93JP-0004125.
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                                                                                                                                                                                                                                                                       94EP-0100276.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation or arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cyclic pentapeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
Les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                       Endo S, Inooka H,
                                                       Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 AA;
                                                                                               Misc-difference
                  Modified-site
                                                                                                                                                                                                                                                                       11-JAN-1994;
                                                                                                                                                                                                                                                                                                           13-JAN-1993;
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                                                                                                                                                                                             EP606881-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This peptide is one of 14 endothelin derivatives which are preferred for use in conjugates with novel bis-(N' (mercapto-ethyl-or -propyl) aminocarbonylmethyl) oxide, sulphide or selenide compounds. The chalcogen-containing compounds are ligands which are land that a technetium or rhenium radioisotope. The iligand-peptide conjugates are useful for non-invasive, in vivo imaging, e.g. of receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bis-(N-(mercapto-ethyl- or -propyl) amino:carbonyl:methyl) oxide, sulphide or selenide and derivs. - are ligands for radioactive technetium or rhenium for diagnosis and therapy, e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-Glu, and this amino acid is part of a cyclic structure formed by condensation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .
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                              | ...|
| | ...|
| note= "attached to chalcogen-contg. ligand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   non-invasive in-vivo visualisation of receptors
                                                                                                                                                                                                                                                                                                                                                                                                         (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                  'note= "D-form residue"
                                                                                                                                                       /label= alle
/note= "D-form residue"
                                                                                                                                                                                                                /note= "D-form residue"
                                                                               see comments"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                              Hilger C, Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR69266 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                     93DE-4311022.
                                                                                                                                                                                                                                                                                                                                93DE-4311022.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 AA;
                                                                                               Misc-difference
                                                                                                                                                                                             Misc-difference
                                                                                                                                      Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                              Dinkelborg L,
                                                                                                                                                                                                                                                                                                                                31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                    Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 38
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New cyclic pentapeptides are disclosed which have a gamma-turn and a beta-turn and which are of formula Cyclo(-Al-A2-A3-A4-A5-) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Bither D-amino acid residues are chosen for Al, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and D-residues for A2 and A5. The peptides are chosen for antagonists against the NKZ receptor or endothalin receptor. They can be used as antisathmatic, antiinflammatory or antiarthritic can be used as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New cyclic penta:peptide(s) having a gamma-turn and a beta-turn used as NK2 receptor antagonists for treating asthma,
/note= "D-Trp; and this amino acid is condensed onto the N-terminal to form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            present sequence is a specifically claimed example of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiasthmatīc; antiinflammatory; antiarthritic; affinity purification; cyclic.
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                       Example 1; Page 9; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR69270 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                Endo S, Inooka H, Kikuchi T,
                                                                   'note= "D-Phe"
                                                                                                                                                                               94EP-0100276.
                                                                                                                                                                                                     93JP-0004125
                                                                                                                                                                                                                          (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-MAY-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                      WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                        Misc-difference 3
                                                                              Misc-difference 5
   Key
Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pentapeptides.
                                                                                                                                                                              11-JAN-1994;
                                                                                                                                                                                                    13-JAN-1993;
                                                                                                                                                          20-JUL-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2;
                                                                                                                                    EP606881-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR69270;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR69270
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                                                                                                                                                                                                                                                                                                                                                     New cyclic pentapeptides are disclosed which have a gamma-turn and a beta-turn and which are of formula Cyclo(-A1-A2-A3-A4-A5-) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Bither D-amino acid residues are chosen for A1, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and antagonists against the NK2 receptor or endothelin receptor. They can be used as antiasthmatic, antiinflammatory or antiarthritic agents or as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                       New cyclic penta:peptide(s) having a gamma-turn and a beta-turn used as NK2 receptor antagonists for treating asthma, inflammation or arthritis
/note= "D-Glu; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
                                                                                     /note= "D-Trp; and this amino acid is condensed
onto the N-terminal to form a cyclic
peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
antiasthmatic; antiinflammatory; antiarthritic;
affinity purification; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a specifically claimed example of the cyclic pentapeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                             Wakimasu M;
                                           "Thr (Bzl) "
                                                                                                                                                                                                                                                                                                                                    Claim 11; Page 16; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAR69268 standard; peptide; 5 AA
                                                                                                                                                                                                                                             Inooka H, Kikuchi T,
                                                                 'note= "D-Leu"
                                                                                                                                                                                                                      (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                             94EP-0100276
                                                                                                                                                                                                  93JP-0004125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-MAY-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2; Conservative
                                             note=
                                                                                                                                                                                                                                                                  WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                    Misc-difference 3
                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 A.
                                 Modified-site
                                                                                                                                                                             11-JAN-1994;
                                                                                                                                                                                                 13-JAN-1993;
                                                                                                                                                       20-JUL-1994.
                                                                                                                                 EP606881-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR69268;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                             Endo S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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beta-turn and which are of formula Cyclo(.A1-A2-A3-A4-A5.) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Either D-amino acid residues are chosen for A1, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and antagonists against the NC2 receptor or endothelin receptor. They can be used as antiasthmatic, antiinflammatory or antiathritic agents or as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cyclic pentapeptides are disclosed which have a gamma-turn and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New cyclic penta:peptide(s) having a gamma-turn and a beta-turn
used as NK2 receptor antagonists for treating asthma,
inflammation or arthritis
/note= "D-Glu; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
antiasthmatic; antiinflammatory; antiarthritic;
affinity purification; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15; DB 15; Length 5; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a specific example of the cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               65.2%; Scort
100.0%; Pred. No. '...
'... 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wakimasu M;
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AAR65728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                             New cyclic penta:peptide(s) having a gamma-turn and a beta-turn used as NK2 receptor antagonists for treating asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                   /note= "D-Asp; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
                                                                                  /note= "D-Trp; and this amino acid is condensed
  onto the N-terminal to form a cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chalcogen; ligand; complex; endothelin; non-invasive imaging;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   technetium, rhenium, in vivo imaging, atherosclerotic plaque, bifunctional chalcogen-interrupted chelate former, cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a specific example of the cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ilarity 100.0%; Pred. No. 7.8
Conservative 0; Mismatches
                                                                                                                                                                                                                                     Endo S, Inooka H, Kikuchi T, Wakimasu M;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                         Example 5; Page 15; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR65742 standard; peptide; 5 AA.
                                                                                                        peptide"
                                                             note= "D-Leu"
                                                                                                                                                                                            93JP-0004125.
                                                                                                                                                                                                                (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                        94EP-0100276.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-JUN-1995 (first entry)
                                                                                                                                                                                                                                                                                                    inflammation or arthritis
                                                                                                                                                                                                                                                           WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 2; Conserva
                                                   Misc-difference 3
 Key
Misc-difference 1
                                                                          Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pentapeptides.
                                                                                                                                                                      11-JAN-1994;
                                                                                                                                                                                            13-JAN-1993;
                                                                                                                              EP606881-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR65742;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 42
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                         5
/note= "the amino group of D-Glu forms a peptide
linkage with the carboxy group of D-Trp"
                                                                                                                                                                                                                                                                                                                                       /note= "the amino group of D-Glu forms a peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bifunctional chalcogen-atom interrupted chelate formers - for radioactive isotopes, their prodn. and use in diagnostics and therapy, e.g. for non-invasive in-vivo representation of receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chalcogen; ligand; complex; endothelin; non-invasive imaging; technetium; rhenium; in vivo imaging; atherosclerotic plaque; bifunctional chalcogen-interrupted chelate former; cyclic.
                                                    label= cyclic
note= "attached to chalcogen-contg. ligand,
                                                                                                                                                                                                                                                                                                                                                                   linkage with the carboxy group of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dinkelborg L, Hilger C, Kramp W, Schier H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred ...
                                                                                                                                                                                                           'label= aIle
'note= "D-form residue"
                                                                                                                                                                                                                                                                                        'note= "D-form residue"
                                                                                                                                                       /note= "D-form residue"
                                                                                                        see comments"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR65728 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 9; Page 11; 13pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93DE-4311023.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93DE-4311023,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.0
les 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUN-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      atherosclerotic plaques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1994-311305/39.
                                                                                                                                                                                                                                                               Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA;
                                                                                                                                 Misc-difference
                                                                                                                                                                                     Modified-site
                                                                                                                                                                                                                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                              Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-OCT-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DE4311023-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 LW
     Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 43
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WPI; 1994-332837/41
Misc-difference 1
                                                                                                                                                                                                                                                                                                                                        5 AA;
                 Misc-difference
                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
Modified-site
                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-AUG-1995
                                                                                                               29-MAR-1994;
                                                                              WO9422491-A
                                                                                               13-0CT-1994
                                                                                                                                                                                                                                                                                                                                                                                           4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR64367;
                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                          4 LW
                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 45
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                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                    preferred for use in conjugates with novel, bifuctional chalcogen-
interrupted chelate forming compounds. The chalcogen-containing
compounds are ligands which are complexed with a technetium or
rhenium radioisotope. The ligand-peptide conjugates are useful
for non-invasive, in vivo imaging, e.g. of receptors or of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide; ligand; radioactive; peptide complex; imaging; receptor; steroid; hormone; atherosclerotic plaque; targetting; endothelin; analogue; derivative; antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                     .
0
                          1..5
/label= cyclic
/note= "attached to chalcogen-contg. ligand,
see comments"
                                                                                                                                                                                                                                                                                                            This peptide is one of 14 endothelin derivatives which are
                                                                                                                                                                                                                                                       Bifunctional, chalcogen-interrupted chelate former for radioisotopes - used in diagnostics and therapy, e.g. for non-invasive in-vivo representation of receptors
                                                                                                                                                                                                                                                                                                                                                                                                 65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; Ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "Forms peptide linkage with DTrp5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cyclic peptide ligand #2 for radiopharmaceutical agent.
                                                                                                                                                                                                   (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                    note= "D-form residue'
                                                                                             'note= "D-form residue'
                                                                                                               /note= "D-form residue"
                location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                      Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR63183 standard; peptide; 5 AA
                                                                                                                                                                                                                                                                                          Claim 9; Page 12; 13pp; German.
                                                                                    /label= alle
                                                                                                                                                                  93DE-4310999
                                                                                                                                                                                    93DE-4310999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-JUN-1995 (first entry)
                                                                                                                                                                                                                      Dinkelborg L, Hilger C,
                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                atherosclerotic plaques.
                                                                                                                                                                                                                                       WPI; 1994-311297/39.
                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                           Misc-difference
                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                  S AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Modified-site
                                                                                                                                                                   31-MAR-1993;
                                                                                                                                                                                   31-MAR-1993;
                                                                                                                                DE4310999-A
                                                                                                                                                06-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                        Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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The sequences given in AAR73170-83 represent peptide ligands which are used with the radioactive peptide complex of the invention. The complexes may be used to image receptors, specifically of steroid hormones, receptor contg. issue and/or atherosclerotic plaque. These peptides allow the complex to be targetted to a specific tissue. These peptides are pref. derived from endothelin, and represent analogues, derivatives or antagonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "optionally has a carboxyl, amido, hydrophobic or macromolecular carrier gp. attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "optionally has an amino, acetyl, 9-fluorenylmethoxy-carbonyl, hydrophobic or macromolecular carrier gp. attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New radio-pharmaceutical agents deriv. from peptide chelating ligands - also the new ligands and complexes with tissue targetting peptide(s), useful for imaging and treatment of e.g. atherosclerosis or steroid dependent cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DP-178 homologue 36 derived from HIV-1 has antiviral activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antiviral activity; DP-178; DP-107; diagnostic; HIV-1LAI; human immunodeficiency virus; transmembrane protein; gp41; alpha helix; leucine zipper; DP-185.
                                                                                                                                                                                                 /note= "Forms peptide linkage with DGlu1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schulze P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rohlfs G,
'note= "D-form residue"
                                                                                                                                /note= "D-form residue"
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                                                            /note= "alloDIle"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dinkelborg L, Erber S, Noll B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR64367 standard; Peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 66; 81pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                            94WO-DE00369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93DE-4311021
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les 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5
/note= "
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/note= "D or L form residues"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                      AAR59460 standard; peptide; 5 AA.
                                                                                                                                                                                                        Claim 1; Page 2; 7pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93JP-0109456.
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                                                                                93JP-0109456
                                                                                                     93JP-0109456
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                             WPI; 1995-041168/06.
                                                                                                                                                                                                                                                                                               Local Similarity
les 2; Conserv
                                                                                                                        (KAOS ) KAO CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KAOS ) KAO CORP
                                                                                                                                                                                                                                                                     5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
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          Modified-site
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Modified-site
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                                         JP06321755-A.
                                                                                 11-MAY-1993;
                                                                                                     11-MAY-1993;
                                                                                                                                                                                     pigmentation
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                                                            22-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                               4 LW 5
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                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                           AAR59460;
                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                         truncated at the amino terminus. DP-178 corresponds to amino acids 638 to 673 of the HTV-1 isolate LAI transmembrane protein 1991. It ectodomain, and complexes with DP-107 (corresponds to amino acids 558-595) which contains a leucine zipper motiff. The peptides complex activity. The peptide derivs. were identified by a computer assisted peptide sequence search. The peptides inhibit transmission to uninfected cells, and can also be used as type and/or subtype uninfected cells, and can also be used as type and/or subtype
                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                  AAR64365-97 are peptide derivs. of DP-178 (AAR64364) which have been
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "forms peptide bond with Xaa at position 5" /note= "D or L form residues"
                                                                                                                                                               Computer search generated synthetic peptides - are inhibitors of
                                                                                                                                                                                                                                                                                                                                                                                    ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Melanin pigmentation inhibition; skin whitening cosmetic; cyclic peptapeptide; sunburn fleck formation treatment.
                                                                                                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 16; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                               Barney SO, Bolognesi DP, Lambert DM, Matthews TJ;
Petteway SR, Wild CT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Melanin pigmentation inhibiting cyclic pentapeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label= Glu, Asp
/note= "D or L form residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /label= Ala, Pro, Hyp
/note= "D or L form residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'note= "D or L form residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                              Claim 11; Page 133; 182pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR59459 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /label= Val, Asp
                                                   94WO-US05739.
                                                                      93US-0073028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-SEP-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 2; Conserva
                                                                                                                                            WPI; 1995-036105/05.
                                                                                          (UYDU-) UNIV DUKE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference 5
                                                                                                                                                                          HIV transmission
                                                                                                                                                                                                                                                                                                                                         5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified-site
                                                07-JUN-1994;
                                                                      07-JUN-1993;
         WO9428920-A
                              22-DEC-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                      4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                         1 LW 2
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                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 46
qq
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "forms peptide bond with Leu at position 1"
/note= "forms peptide bond with Leu at position 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR59459-R59461 are cyclic pentapeptides which inhibit melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Melanin pigmentation inhibition; skin whitening cosmetic; cyclic peptapeptide; sunburn fleck formation treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         65.2%; Score 15; DB 16; Length 5; llarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
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Gaps

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The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                       /label= OTHER
/note= "alpha-amino acid with conformationally
restricted 4-16C side chain, preferably
1,2,3,4-tetrahydro-2-carboline-3-carboxylic
acid"
                                                                                                                                                                                                                                                                           Endothelin receptor antagonist, organ hypofunction, liver; surgery; transplant, cyclic.
                                                                                                                                                                                                                                                  Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                                                                                    /note= "D-Asp, which is part of a cyclic peptide formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "D-Trp, which is part of a cyclic per
formed by condensation with D-Asp(1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
65.2%; Score 15; DB 16; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
                             Length 5;
                                                     Indels
                          65.2%; Score 15; DB 16; I
100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 34; Page 106; 107pp; English.
                                                                                                                                                                     AAR69429 standard; peptide; 5 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94EP-0106093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93JP-0094332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                          (first entry)
                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wakimasu M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-000740/01.
                                     Local Similarity
les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Misc-difference 3
    5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-APR-1993;
                                                                                                                                                                                                                       18-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EP626174-A.
                                                                              4 LW 5
                                                                                                      1 LW 2
                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kitada C,
                                                                                                                                                                                                AAR69429;
     Sequence
                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                     Matches
                                                                                                                                             RESULT 49
                                                                                                                                                           AAR69429
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                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "forms peptide bond with Asp at position
                                                                                                                                                                                                                       .;
0
                                                                                                             AAR59459-R59461 are cyclic pentapeptides which inhibit melanin pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR59459-R59461 are cyclic pentapeptides which inhibit melanin pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                                    Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                          Melanin pigmentation inhibition, skin whitening cosmetic, cyclic peptapeptide, sunburn fleck formation treatment.
                                                                                                                                                                                              Length 5;
                                                                                                                                                                                           65.2%; Score 15; DB 16; Length 5; larity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                  Melanin pigmentation inhibiting cyclic pentapeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                      Example 1; Page 3; 7pp; Japanese.
                                                                                                                                                                                                                                                                                                                                     AAR59461 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 3; 7pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93JP-0109456
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                                                                                                                                                                                                                                                                                                                                                                                        01-SEP-1995 (first entry)
                                                                                                                                                                                                       Local Similarity
             WPI; 1995-041168/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1995-041168/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      KAOS ) KAO CORP.
                                                                                                                                                                    5 A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modified-site
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                                                              pigmentation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pigmentation
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                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                AAR59461;
                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                      Matches
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AAR69414 standard; peptide; 5 AA.
                                                                (first entry)
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es 2; Conserv
                                                                                                                                    transplant; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA;
                                                                                                                                                                                                       Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                     20-APR-1994;
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                                                                  18-JUL-1995
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                                       AAR69414;
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AAR69414
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                           Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                     Endothelin receptor antagonist; organ hypofunction; liver; surgery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or
                                                                                                                                                                                                                                                                                                                  /note= "D-Ser, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-Trp, which is part of a cyclic peptide
formed by condensation with D-Ser(1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .,
                                                                                                                                                                                                                                                                                                                                                                      /note= "alpha-amino acid with conformationally
    restricted side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 16; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 34; Page 106; 107pp; English.
                                                                                                           AAR69430 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                               note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                           /label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94EP-0106093.
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                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transplant of the organ
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Best Local Similarity
Matches 2; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1995-000740/01.
                                                                                                                                                                                                                                    transplant; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 3
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                                                                                                                                                                18-JUL-1995
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                                                                                                                                      AAR69430;
              4 LW
                                                                              RESULT 50
AAR69430
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/note= "D-Cys(O3S), which is part of a cyclic peptide formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Endothelin receptor antagonist; organ hypofunction; liver; surgery; transplant; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                   Endothelin receptor antagonist; organ hypofunction; liver; surgery;
Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                              /note= "D-Trp, which is part of a cyclic peptide
formed by condensation with D-Cys(1)"
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                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR69415 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                          /note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94EP-0106093.
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RESULT 51

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The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endothelin receptor antagonist; organ hypofunction; liver; surgery;
transplant; cyclic.
                              /note= "D-Trp, which is part of a cyclic peptide formed by condensation with D-Asp(1)"
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                                                                                                                                                                                                                                                                                                        Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 16; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "Tyr-NH-Ind-OH, where Ind is
1-carboxyindane-2-y1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "hexamethyleneimino-CO-Leu"
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                                                                                                                                                                                                                                             Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                          Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR69392 standard; peptide; 5 AA.
/note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'note= "D-Ala"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'label= bAla
                                                                                                                                             94EP-0106093.
                                                                                                                                                                             93JP-0094332
                                                                                                                                                                                                             (TAKE ) TAKEDA CHEM IND LID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 100.00
Best Local 2; Conservative
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                                                                                                                                                                                                                                            Wakimasu M,
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                  Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 AA;
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                                                                                                                                             20-APR-1994;
                                                                                                                                                                             21-APR-1993;
                                                                                                               30-NOV-1994.
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                                                                               EP626174-A
                                                                                                                                                                                                                                            Kitada C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin receptor antagonist; organ hypofunction; liver; surgery;
                                                                                                                                   /note= "D-Thg(2) or D-Cpg, where Thg(2) is 2-(2-thienyl)glycine and Cpg is 2-cyclopentyl-glycine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or transplant of the organ
                                                                           /note= "D-Asp, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                        /note= "D-Trp, which is part of a cyclic peptide formed by condensation with D-Asp(1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "D-Asp, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
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                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                        Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR69413 standard; peptide; 5 AA.
                                                                                                                           /label= OTHER
                                                                                                                                                                                                                                                                                                                        94EP-0106093
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    Kitada C, Wakimasu M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-000740/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                       Misc-difference 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 AA;
                                                           Misc-difference
                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                      20-APR-1994;
                                                                                                                                                                                                                                                                                                                                                      21-APR-1993;
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                                                                                                                                                                                                                                                         EP626174-A
             Synthetic.
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Synthetic.
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                                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 56
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                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      part of a cyclic structure obtained by condensation of D-Trp(5) onto the alpha-NH2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   part of a cyclic structure formed by condensation of the COOH onto the NH2 group of D-Glu(1)"
                                                                                                                                                                                                                                                                                                                                                                                                     diagnostic imaging, atherosclerotic plaque; tumour; inflammation; conjugate; radiotherapy; cysteine-free; metal-binding; endothelin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "D-form residue; and this amino acid forms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-form residue; and this amino acid forms
                                                                                                Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or
                                                                                                                                                                                                                                                                                                                                                                                   Endothelin sequence for use as conjugate with Cys-free peptide.
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                                                                                                                                                                                                                             65.2%; Score 15; DB 16; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                               Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                    Claim 7; Page 83; 107pp; English.
                                                                                                                                                                                                                                                                                                                               AAR86453 standard; peptide; 5 AA.
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          94EP-0106093.
                                            (TAKE ) TAKEDA CHEM IND LTD.
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                           93JP-0094332.
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                                                                                                                                                                                                                                               Conservative
                                                                                                                    transplant of the organ
                                                                                                                                                                                                                                     Local Similarity
les 2; Conserval
                                                                                WPI; 1995-000740/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 5
                                                                                                                                                                                                            5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
          20-APR-1994;
                           21-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DE4337599-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                 4 LW 5
                                                                                                                                                                                                                                                                                    1 LW 2
                                                                                                                                                                                                                                                                                                                                                 AAR86453;
                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                        cyclic.
                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                              RESULT 55
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conjugates with peptides, proteins, biomolecules and macromolecules, conjugates with peptides, proteins, biomolecules and macromolecules, their complexes with metal ions and their water-soluble salts; where their complexes with metal ions and their water-soluble salts; where including at least one Met, Arg, Lys or Asn but no Cys; R1 is H or a hydrocarbyl or acyl group, or a bond to a conjugate; and R2 is OH, hydrocarbyloxy or an optionally substituted amine group (forming an amide), or a bond to a conjugate. The peptides are useful when complexed to a metal (e.g. 99m-TC) for in-vivo diagnostic imaging and therapy of tumours, organs and fooi of inflammation. Conjugates of the peptides can be targetted to specific tissues. A particular application is imaging of atherosclerotic plaque. Metal is bound with good in-vivo stability; unbound material is rapidly cleared; and there is no allergic reaction. Since the peptides are free of Cys, they can be prepared simply and rapidly and can be presented in the form of a kit for radio-pharmaceutical production. Specific examples of the peptides are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sex-hormone, human; veterinary medicine; cancer; prostate; uterus; breast; pituitary; prostatomegaly; endometriosis; hysteromyoma; precocious puberty; amenorrhoea; premenstrual syndrome; cyclic; multilocular ovarian syndrome; comedo; pregnancy; contraception; ovulation; menstrual cycle; cestrus; meat quality; growth; spawning; fish; testosterone; superagonist; leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "N-pToluenesulphonylarginine, D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pharmaceutical production. Specific examples of the peptides are given in AARB6424 - AARB6439.
The present sequence is a specific example of an endothelin peptide which can be used as a conjugate with the new peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Luteinising hormone releasing hormone receptor; LH-RH; antagonist;
Frommel C, Hohne W;
ier HM, Schneider-Mergener J;
    Conrad J, Dinkelborg L, Erber S, Frommel C, Hohne W;
Kramp W, Kuttner G, Malin R, Schier HM, Schneider-Mergener Steinbrecher R, Froemmel C, Hoehne W, Kuettner G, Schler H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Joined via a peptide linkage to Phel"
                                                                                                                                                                                                                                                                 New cysteine-free peptide(s) and their metal ion complexes conjugated to targetting agent, useful for in vivo imaging of tumours, atherosclerotic plaque etc. and for radiotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cyclic pentapeptide #25 used in LH-RH receptor antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 16; L larity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW17809 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 13; Page 20; 25pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUL-1997 (first entry)
                                                                                                                                                                                     WPI; 1995-171400/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin; receptor; leukocyte; vascular wall; endothelium; extravasation; inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.
                                                                                                                                                                                                                                     inteinising hormone releasing hormone (LHT.RH) receptor antagonist composition of the invention. Peptides such as this are used to prevent or treat sex-hormone dependent disorders in human or veterinary medicine, esp. cancers (of the prostate, uterus, breast or pituitary), prostatomegaly, endometriosis, hysteromyoma or precocious puberty, but also amenorrhoes, premental syndrome, multilocular ovarian syndrome, also to induce ovulation) and the menstrual cycle. They are also used or control oestrus in animals, to improve meat quality and control growth, and to promote spawning in fish. They may also inhibit the transient increase in testoseerone blood levels caused by admin. of superagonists such as leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                              This peptide represents a cyclic peptide which is included in the
                                                                                                                                                          Cyclic penta:peptide(s), some new, as LH-RH receptor antagonists used to treat or prevent sex hormone dependent disorders, e.g. cancer, also for control of pregnancy and menstruation and to improve meat quality in animals
                                                                                                                                                                                                                                                                                                                                                                                                                .
0
                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 17; Length 5;
Pred. No. 7.8e+05;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "amidated C-terminus"
                                                                                                                                                                                                          Example 45; Page 173; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Anti-ELAM-1 binding peptide #213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR86236 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "Ac-Leu"
                                                                                                                    Furuya S, Kato K, Kitada C;
                                                                                                                                                                                                                                                                                                                                                                                         65.2%;
                                                                  95JP-0110933.
                                                  96WO-JP01140
                                                                                                 (TAKE ) TAKEDA CHEM IND LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JUN-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                              2; Conservative
                                                                                                                                       WPI; 1996-497569/49.
                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Key
Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Modified-site
                                               5-APR-1996;
          WO9634012-A1
                                                                  09-MAY-1995;
                                                                              28-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                1 FXXLW 5
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                              31-0CT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR86236;
                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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that bind to endothelial leukocyte adhesion molecule (BLAM)-1. This that bind to endothelial leukocyte adhesion molecule (BLAM)-1. This molecule is a member of the selectin family of receptors and is involved in binding of leukocytes to the vascular endothelial wall prior to extravasation of the leukocyte, e.g. to a site of inflammation.

The peptides bind pref to B-selectin but may also bind L- or P-selectin, and can be used to treat conditions mediated by E-selectin, e.g. inflammatory conditions. The peptides have strong affinity for the selectin receptors and inhibit the binding of the sialyl Lewis (SLE-x) car of cell surface glycoproteins to B-selectin. The peptide are small, generally less than 2 kD, have an IC50 of up to 100 micromole against binding of HL60 cells to BLAM-1, have one or more peptide linkages replaced by CH2OC(O)NR, phosphonate, CH2SOZNR, CH3ONR, CON(R6), or NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl. The peptides may also have substituted N- and C-termini e.g. cucinimido, N-benzyloxycarbonyl or N-lower alkyl cpds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                 New peptide(s) that bind to endothelial leukocyte adhesion molecule 1 - useful for treating inflammation and other E-selectin mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bifunctional, bis-nicotinamide, chelator, radioactive, technetium, rhenium, conjugate, endothelin, angiotensin, targetting agent, organ, in vivo imaging, receptor; tissue, atherosclerotic plaque; diagnosis, breast, prostatic; carcinoma; cytostatic agent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Targetting conjugate for bis-nicotinamide bifunctional chelator.
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                                                                                                                                                                                            Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;
Martens CL, Ruhland-fritsch B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 17; L 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'note= "allo-D-form residue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 73; 85pp; English.
                                                                                                                                  (AFFY-) AFFYMAX TECHNOLOGIES NV
                                                                        94US-0241054.
      95WO-US06315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-FEB-1998 (first entry)
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                                                                                                                                                                                                                                                                                                   WPI; 1996-010687/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  delivery; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
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11-MAY-1995;
                                                                        11-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 LW 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                               diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW33328
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This peptide is an endothelin antagonist which is the active component in a new cerebral oedema inhibitor, used to prevent swelling in head injury, cerebral tumour, blood vessel damage in apoplexy, hypertension, respiratory failure, carbon monoxide poisoning, hyponatraemia, acute kidney damage, hyperglycaemia, hypoglycaemia, adrenal function failure, collagen disease, inadequate buffering of blood, or poisoning by tin, lead or arsenic. The inhibitor prevents cerebral hernia, headache, nausea, dizziness, restlessness and loss of consciousness, especially sudden death from cerebral hernia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bifunctional sulphonamide; ligand; radioactive; technetium; rhenium; conjugate; endothelin; angiotensin; targetting agent; organ; in vivo imaging; receptor; tissue; atherosclerotic plaque; diagnosis; breast; prostatic; carcinoma; cytostatic agent;
                                                                                                                                                                         Cerebral oedema inhibitor comprising endothelin antagonist - esp. specified oleanonic acid deriv., cyclopeptide or bosentan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Targetting conjugate for bifunctional sulphonamide type ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                           Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 18; L ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "allo D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          'note= "D-form residue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW33020 standard; peptide; 5 AA.
                                                                                                                                                                                                                   Claim 2; Page 12; 20pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95DE-1036785.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      95DE-1036785.
                          96WO-JP01438.
                                                        95JP-0160060.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-JAN-1998 (first entry)
                                                                                      (SHIO ) SHIONOGI & CO LTD.
                                                                                                                  Matsuo Y, Ninomiya M;
                                                                                                                                              WPI; 1997-034102/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
es 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         delivery; cyclic
                                                                                                                                                                                                                                                                                                                                                                                                    5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DE19536785-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-SEP-1995;
                             29-MAY-1996;
                                                        02-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-1997
05-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW33020;
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 60
δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   QD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "D-form residue. This Trp is not a true C-terminal residue but is part of a cyclic peptide.
The alpha carboxyl group is condensed with the alpha amino group of Asp (1)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "D-form residue. This Asp is not a true N-terminal residue but is part of a cyclic peptide.

The alpha amino group is condensed with the alpha carboxyl group of Trp(5)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cerebral hernia; tumour; apoplexy; hypertension; respiratory failure; hyperglycaemia; hypoglycaemia; adrenal function failure; collagen disease; poisoning; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                           Novel bis-nicotinamide bifunctional chelating agent for radioactive technetium and rhennium, when conjugated to an endothelin or angiocensin derived targetting agent, e.g. the present peptide, can be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used for cytostatic agent delivery.
                                                                                                                                                                                                                        New bis-nicotinamide bi:functional chelating agents - forming new, stable technetium or rhenium complexes and conjugates useful as radiodiagnostic and radio:therapeutic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Endothelin antagonist #1 used as brain oedema inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Endothelin antagonist; brain oedema inhibitor; bosentan;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 18; L
llarity 100.0%; Pred. No. 7.8e+05;
Conservative 0; Mismatches 0;
                                                                                                                                                 Hilger CS, Kramp W, Platzek J;
                                                                                                                     (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW32229 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                   Claim 11; Page 42; 47pp; German.
                                                           96WO-DE01824.
                                                                                         95DE-4036783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-FEB-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                             WPI; 1997-212562/19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                   5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified-site
                                                                                                                                                 Dinkelborg L,
Raduechel B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modified-site
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   WO9710853-A2
                                                             19-SEP-1996;
                                                                                         21-SEP-1995;
                                27-MAR-1997.
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4 LW 5
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                                                                                                                                                                                                                               RESULT 62
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                                                          Bifunctional sulphonamide type ligands for radioactive technetium or rhenium - and their chelates and conjugates, useful for imaging
                                                                                                                                                                                                                       Gaps
                                                                                                                                 can
                                                                                                                      technetium and rhenium, when conjugated to an endothelin or angiotensin derived targetting agent, e.g. the present peptide, ca be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used for cytostatic agent delivery.
                                                                                                                                                                                                                                                                                                                                                                         rhenium; conjugate; endothelin; angiotensin; targetting agent; organ; in vivo imaging; receptor; tissue; atheroselerotic plaque; diagnosis; breast; prostatic; carcinoma; cytostatic agent; delivery; cyclic.
                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                    Targetting conjugate for bifunctional sulphonamide type ligand
                                                                                                              Novel bifunctional sulphonamide type ligands for radioactive
                                                                                                                                                                                                                                                                                                                                                                   Bifunctional sulphonamide, ligand, radioactive, technetium,
                                                                                                                                                                                                   65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; rative 0; Mismatches 0; Indels
                  Platzek J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Platzek J;
                  Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Kramp W,
(UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "allo D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note= "D-form residue'
                  Hilger CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hilger CS,
                                                                                                                                                                                                                                                                                                AAW11213 standard; peptide; 5 AA
                                                                                              Claim 9; Page 12; 12pp; German.
                                                                            carcinoma or atherosclerosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95DE-1036780
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                                                                                                                                                                                                                                                                                                                                    26-JAN-1998 (first entry)
                                                                                                                                                                                                                     Conservative
                  Erber S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Erber S,
                                          WPI; 1997-193894/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-193892/18.
                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference 1
                                                                                                                                                                                  5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                 Dinkelborg L,
Raduechel B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dinkelborg L,
Raduechel B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DE19536780-A1
                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21-SEP-1995;
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                                                                                                                                                                                                                                     LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                      ß
                                                                                                                                                                                                                                                                                                                 AAW11213;
                                                                                                                                                                                   Sequence
                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                      3
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                                                                                                                                                                                                                     Matches
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Gaps
                                                                                                                                                                                                      technetium and rhenium, when conjugated to an endothelin or angiotensin derived targetting agent, e.g. the present peptide, ca be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used for cytostatic agent delivery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bifunctional sulphonamide type ligands for radioactive technetium and rhenium - and their chelates and conjugates with targetting agents, useful for imaging carcinoma, atherosclerotic plaque, etc.
Bifunctional sulphonamide type ligands for radioactive technetium and rhenium - and their chelates and conjugates with targetting agents, useful for imaging carcinoma, atherosclerotic plague, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rhenium; conjugate; endothelin; angiotensin; targetting agent; organ; in vivo imaging; receptor; tissue, atherosclerotic plaque; diagnosis; breast; prostatic; carcinoma; cytostatic agent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Targetting conjugate for bifunctional sulphonamide type ligand.
                                                                                                                                                                              Novel bifunctional sulphonamide type ligands for radioactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bifunctional sulphonamide; ligand; radioactive; technetium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15; DB 18; Length 5;
Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Platzek J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         65.2%; Scor.
100.0%; Pred. No. ...
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Erber S, Hilger CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW11212 standard; peptide; 5 AA.
                                                                                                                      Claim 11; Page 19; 19pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 11; Page 19; 19pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "D-form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95DE-1036780
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1997-193892/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 delivery; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
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Raduechel B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DE19536780-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New sulphide-sulphonamide bi:functional chelating agents - forming new, stable technetium or rhenium complexes and conjugates useful as radiodiagnostic and radio-therapeutic agents
                            technetium and rhenium, when conjugated to an endothelin or angiotensin derived targetting agent, e.g. the present peptide, can be used for the in vivo imaging of corgans, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      XSNS-type bifunctional sulphide containing sulphonamide chelating agents and their technetium or rhenium complexes, can form covalently bonded conjugates with subserses selectively enriched in diseased tissue, e.g. the present peptide. The substance is an endothelin or angiotensin peptide, or a partial sequence, analogue, derivative or antagonist, or a chemotactic peptide. The conjugates are radiodiagnostic or radiotherapeutic agents, useful in the diagnosis of tumours, ischaemia or atherosclerotic
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conjugating cyclic peptide for radio-therapeutic/diagnostic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conjugate; diseased tissue; endothelin; angiotensin; chemotactic; radiodiagnostic; radiotherapeutic; diagnosis; tumour; ischaemia; atherosclerosis; vascular disorder; ischemia; circular; cyclic.
               Novel bifunctional sulphonamide type ligands for radioactive
                                                                                                                                                                         Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Platzek J;
                                                                                                                                                                                                       0; Indels
                                                                                                                                                                       65.2%; Score 15; DB 18; L
100.0%; Pred. No. 7.8e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hilger CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                 AAW14986 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 13; Page 45; 50pp; German.
                                                                                                             for cytostatic agent delivery.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95DE-4036781.
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                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Erber S,
                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-202627/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference 3
                                                                                                                                           5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dinkelborg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                   4 LW 5
                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                             AAW14986;
                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                  63
                                                                                                                                                                                                                                                                                                                 RESULT
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                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Peptide binding to low density lipoprotein - useful in removing LDL from body fluids and as carrier peptide(s) for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
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0
                                                                          ;
                                                                                                                                                                                                                                                                                              density; lipoprotein; LDL; binding; absorbent; removal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; Live 0; Mismatches 0; Indels
                                                 Length 5;
                                                                          0; Indels
                                                65.2%; Score 15; DB 18; L
100.0%; Pred. No. 7.8e+05;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                      low density lipoprotein binding peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 11; Page 50; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                      /note= "amidated"
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                                                                                                                                                                                          AAW22915 standard; peptide; 5 AA.
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                                                                          Conservative
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nes 2; Conserva
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                                                                                                                                                                                                                                                                                                            body fluid; reagent
vascular disorders.
                                                         Local Similarity
ses 2; Conserv
                          5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
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                                                                                                                             Low density, lipoprotein; LDL; binding; absorbent; removal; body fluid; reagent.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
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                                                                                      Low density lipoprotein binding peptide
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/note= "amidated"
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(ASAH ) ASAHI MEDICAL CO LTD.
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                                                      02-OCT-1997 (first entry)
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Modified-site
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                   AAW22920;
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Human; beta-amyloid precursor protein; beta-APP; diagnosis; cancer; frameshift mutation; age-related disease; neurodegenerative disorder; Alzehmer, s disease; Down's syndrome; myotonic dystrophy; neuronal; Huntington's disease; multiple sclerosis; alcoholic liver disease; disease; multiple sclerosis; alcoholic liver disease; diabetes mellitus type II; microtubule associated protein; Tau; Big Tau; ubiquitin B; apolipoprotein E; MAP2; neurofilament-L; neurofilament-C; presenilin I; presenilin II; cellular tumour antigen; glial fibrillary acidic protein; GFAP; p53; semaphorin III; HUPF-I; bcl-2; B-cell leukemia/lymphoma 2 proto-oncogene; HMGP-C; NSP-A; high mobility group protein-C; neuroendocrine specific protein A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Peptide binding to low density lipoprotein - useful in removing LDL from body fluids and as carrier peptide(s) for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
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(ROYA-) ROYAL NETHERLANDS ACAD ARTS & SCI.
(UYRO-) UNIV ROTTERDAM ERASMUS.
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(ASAH ) ASAHI MEDICAL CO LTD.
                                                                     96WO-JP01734.
                                                                                                                                           95JP-0176904.
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                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1997-087320/08
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                                                                     21-JUN-1996;
                                                                                                                                           21-JUN-1995;
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09-JAN-1997.
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This invention describes a novel method for the diagnosis of a disease caused by, or associated with, an RNA molecule that has a frameshift mutation. The method is used to diagnose age-related diseases, especially cancer and a wide range of neurodegenerative disorders (e.g. Alzheimer's disease, Down's syndrome, myotonic dystrophy, Huntington's disease, multiple sclerosis, alcoholic liver disease, diabetes mellius type II and many others listed) or susceptibility to these disorders. The method allows a definitive diagnosis of Alzheimer's disease in living patients, at an early stage. It is based on the observation that disease may be caused by mutations in RNA rather than DNA. The invention describes the used of neuronal system RNA molecules, specifically proteins including poteins Tau and Big Tau, ubiquitin B, apolipoprotein E, microtubule associated protein 2 (MAP2), neurofilament-M, mutofilament-M, neurofilament-M, neurofilament-M, neurofilament-M, protein (GRAP), the cellular tumour antigen p53, B-cell leukemia/lymphoma 2 (bcl-2) prote-oncogene, semaphorin III, HUPF-1, high mobility group protein-C (HMGP-C) and neuroendocrine specific protein A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "D-Glu is condensed onto D-Trp at position 5 to form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tyrosine kinase blocker; chemotherapy; antibody; conjugate; cyclic.
                                                                                                     Diagnosing disease by detecting frameshift mutations in RNA or corresponding protein mutations - used to diagnose cancer and neurological diseases, particularly Alzheimer's disease, and also for treatment and prevention with specific ribozymes or wild-type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   endothelin receptor; antagonist; atherosclerosis; radionuclide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 19; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
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                 Van Leeuwen FW;
                                                                                                                                                                                                                       Disclosure, Figure 6; 258pp; English.
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                 Burbach JPH, Grosveld FG,
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                                                   WPI; 1998-609901/51.
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tes 2; Conserv
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                                                                       N-PSDB; AAX75757
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Matches
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The invention relates to (A) the use of conjugates of the following formula as therapeutic agents: B-Wn; where E = a residue which can bind an endothelin receptor, and is derived from an endothelin partial sequence or an endothelin an endothelin partial sequence or an endothelin an endothelin partial sequence or an endothelin an endothelin derivative, an endothelin partial sequence or an endothelin antagonist; W = an active group which: (i) is a radionuclide or conflormed an endothelin antigony an antibody, an antibody fragment, a complex with a conflormed ragery an antibody, an antibody fragment, a peptide, a carbohydrate, an endicament, a thrombocyte aggregation inhibitor, an anti-inflammatory, a calcium antagonist, a lipid lowering agent or anti-inflammatory a calcium antagonist, a lipid lowering agent or anti-inflammatory a calcium antagonist, a lipid lowering agent or an enemotherapeutic agent, an antibody, an antibody fragment, a peptide, a carbohydrate, an oligonucleotide, a protein tyrosine kinase cof the above formula in which wis a matibody, an antibody fragment, a poptide, a carbohydrate, an oligonucleotide, a protein tyrosine kinase blocker, an anti-thrombotic agent, a growth factor inhibitor, an anti-thrombotic agent, a growth factor inhibitor, an anti-thrombotic agent, a lipid lowering agent or an anti-thrombotic agent, a lipid lowering agent or an anti-thrombotic agent, a lipid lowering agent or an anti-thrombotic agent of cardiovascular disorders such as configeration, subarachmotid haemorrhage, preeclampsia, renal disorders, etc. The conjugates become enriched in cells in which endothelin receptors are expressed. Even at low doses a therapeutically effective enrichment of the active agent at low doses a therapeutically effective enrichment of sarpuence represents a specifically claimed endothelin receptor binding expression in the body, reducing side effects. The prinding expecifically claimed endothelin receptor binding expecifically cannel endothelin receptor binding expecifically cannel endo
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                                                                                                                                                                                                                                             Therapeutic use, e.g. in treatment of atherosclerosis, of endothelin conjugates - which comprise residue which can bind endothelin receptor, conjugated to groups such as radionuclides or protein tyrosine kinase blockers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclic peptide which mimics surface feature of endothelin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 5;
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endothelin-1; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                           Dinkelborg L, Hilger C, Speck U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW50857 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                        Claim 3; Page 15; 21pp; German.
                   96DE-1052374.
                                                             96DE-1052374.
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                                                                                                            (SCHD ) SCHERING AG.
                                                                                                                                                                                                        WPI; 1998-363559/32.
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                   04-DEC-1996;
                                                                  04-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW50857
8
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0; Gaps

. 0

/note= "D-Trp is condensed onto D-Glu at position 1 to form a cyclic peptide"

/label= alle
/note= "D-form allo-isoleucine"

Modified-site

Modified-site

DE19652374-A1 10-JUN-1998.

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/note= "D-form residue, amino group of Asp is
                                                                                                                                                                                                             (TEXA-) TEXAS BIOTECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY30995 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                96US-0747137.
                                                                                                                                                                    90US-0628111.
92US-0900623.
94US-0223513.
                                                                                                                                         94US-0410955.
                                                                                                                                                            94US-0410955.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-OCT-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                           /note=
                                                                                                                                                                                                                                 Chan MF;
                                                                                                                                                                                                                                                    WPI; 1998-239252/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                              Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5 AA;
                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-NOV-1996;
                                                                                                                                         10-NOV-1994;
                                                                                                                                                             10-NOV-1994;
                                                                                                                                                                    14-DEC-1990;
18-JUN-1992;
                                                                                                                                                                                           05-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US5945033-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-AUG-1999
                                                                                                  US5736509-A
                                                                                                                     07-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                                                                                                 Balaji VN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ŋ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY30995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4
LW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX30995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                               The invention relates to cyclic peptides that mimic the surface features of the endothelin peptides. The peptides shown in the invention are pentapeptides, hexapeptides and heptapeptides that contain an L-Trp, D-Trp residue or a derivative of D- or L-Trp, such as N-Me-Trp. The peptides modulate the activity of one or more members of the endothelin family of pepides. They are useful for relieving symptoms of and for treating endothelin mediated disorders and diseases. The present sequence represents a cyclic pentapeptide.
                                                                                        /note= "D-form residue, carboxy group of Trp is
condensed onto the amino group of Glu1 to form
a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                     New cyclic penta-peptides which are mimics of C-terminus of natural endothelin-1 - useful for relieving symptoms of and for treating endothelin mediated disorders and diseases
                             /note= "D-form residue, amino group of Glu is
condensed onto the carboxy group of Trp5 to
form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cyclic peptide which mimics surface feature of endothelin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 19; Length 5; 100.0%; Pred. No. 7.8e+05; rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Endothelin; endothelina; endothelinb; ETa; ETb; receptor;
                                                                       /note= "D-form residue'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
1
                                                                                                                                                                                                                                                                                                                                               Example 3; Column 33; 30pp; English
            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW50859 standard; peptide; 5 AA.
                                                                                                                                                                                                                                             (TEXA-) TEXAS BIOTECHNOLOGY CORP.
                                                                                                                                                                                           94US-0410955.
90US-0628111.
92US-0900623.
94US-0223513.
                                                                                                                                                                         94US-0410955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 endothelin-1; cyclic.
                                                                                                                                                                                                                                                                 Chan MF;
                                                                                                                                                                                                                                                                                   WPI; 1998-239252/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                            Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
Modified-site
                      Modified-site
                                                                               Modified-site
                                                                                                                                                                         10-NOV-1994;
                                                                                                                                                                                                                         05-APR-1994;
                                                                                                                                                                                             10-NOV-1994;
                                                                                                                                                                                                     14-DEC-1990;
                                                                                                                                  JS5736509-A
                                                                                                                                                                                                                18-JUN-1992
                                                                                                                                                     07-APR-1998
                                                                                                                                                                                                                                                                 Balaji VN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ŋ
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW50859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW50859
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The invention relates to cyclic peptides that mimic the surface features of the endothelin peptides. The peptides shown in the invention are pentapeptides, hexapeptides and heptapeptides that contain an L-Trp, D-Trp residue or a derivative of D- or L-Trp, such as N-Me-Trp. The peptides modulate the activity of one or more members of the endothelin family of pepides. They are useful for relieving symptoms of and for treating endothelin mediated disorders and diseases. The present sequence represents a cyclic pentapeptide.
                                                                                                                                                                                                       "D-form residue, carboxy group of Trp is condensed onto the amino group of Aspl to form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Non-crosslinked protein particle; diagnostic; therapy; monodisperse; albumin; haemoglobin; nanometer; micrometer; clearance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New cyclic penta-peptides which are mimics of C-terminus of natural endothelin-1 - useful for relieving symptoms of and for treating endothelin mediated disorders and diseases
condensed onto the carboxy group of Trp5 form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Non-crosslinked protein particle peptide 44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Prec. ...
                                                                                                                                                                                                                                                                             a cyclic peptide"
                                                                                                                                  'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Column 33; 30pp; English.
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                                                                                                                                                                                                                                                                     This invention describes a novel aqueous suspension of monodisperse particles on non-crosslinked, non-denatured albumin (50-5000 nm) which is stable against dissolving upon dilution with an alcohol-free aqueous medium. The method involves (a) forming an aqueous solution containing albumin and hemoglobin and (b) treating the aqueous solution with an alcohol to cause the solution to become turbid. The particles are useful as agents for in vivo administration, either of their own administration or as a vehicle for other therapeutic or diagnostic agents. The method permits the formation of albumin and hemoglobin particles in the nanometer and micrometer size range, in a form closer to their natural form than the forms of the prior art. The particles therefore constitute a more closely controlled agent for in vivo administration, with greater ease of clearance from the body after their period of usefulness.

ANY30952-Y31135 represent peptides used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gene characterisation, recombinase, homology clamp, homology motif tag, HMT; genetic manipulation, genetic engineering, gene transcripion, GPCR, drug target, homologous recombination; target identification; TGF-beta; G-protein coupled receptor; transforming growth factor-beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                            Non-crosslinked protein particles for therapeutic and diagnostic use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Acetyl choline (muscarinic) GPCR consensus peptide motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 20; L ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                           Example 22; Column 65-66; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zeng H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY31277 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zarling D,
               94US-0212546.
91US-0641720.
92US-0959560.
                                                               93US-0069831.
96US-0747137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98WO-US26498.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0070734.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-SEP-1999 (first entry)
                                                                                                              (HEMO-) HEMOSPHERE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Rest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PANG-) PANGENE CORP.
                                                                                                                                                                            WPI; 1999-508153/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lehman CW, Pati S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-458689/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
                                                                             12-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9937755-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-1997;
                                            13-OCT-1992;
                                                               01-JUN-1993;
                              15-JAN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2 LW 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY31277;
                                                                                                                                             Yen RCK;
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The invention provides compositions and methods for the evaluation and characterisation of individual and sets of genes in disease states. The composition comprises at least one recombinase and at least two singlestranded targeting polynucleotides which are substantially complementary to each other and each having a consensus homology clamp for a gene family; i.e. a homology motif tag (HWT). The composition is useful in kit form which include the composition as libraries or pools of degenerate casDNA probes along with other reagents such as recombinase etc. The methods and compositions are used for inactivation of a gene family gene conformation inchlumen animal or plant). This is useful for generating decrease or alter the biological activity of one or more genes in a cell of or transgenic nonhuman animal or plant). This is useful for generating animal models of disease, or in the elucidation of gene function and activity. Alternatively, the biological activity of the wild-type gene activity. Alternatively, the biological activity altered to mimic disease states. This includes genetic amnipulation of non-coding gene sequences that affect the transcription of genes, including promoter, repressors, enhancers and transcriptional activity altered to mimic sequences that affect the transcriptional activity of more compositions are useful in functional genes, uncluding promoter, compositions are useful in functional genes, as well as in election of new drug targets HMTs used in homologous recombination wide variety of related genes, potentially resulting in a wide variety of composition of continuing those related to disease states. This may also be done on a cellular level to identify genes involved in cellular level to identify genes involved in cellular level to identify genes involved in collular level to identify genes involved in collular level to identify genes involved in collular level to identify and transforming growth factor-beta (TGF-beta) gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                     New compositions and methods for targeting sequence modifications in related family genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antigenic peptide, streptokinase, streptokinase-specific antibody, thrombolytic activity, thrombolytic therapy, glomerulonephritis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 20; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antigenic peptide of streptokinase.
                                                                                        Disclosure, Fig 1A; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY01560 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98WO-US17114.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-0055911.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEHO ) GEN HOSPITAL CORP. (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUN-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Streptococcus equisimilis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 2; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rheumatic fever.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9908698-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       35-FEB-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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5 AA;
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                                                                                                                        4 LW 5
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                                                                                                                                                                                                                                                                               AAB52657;
                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Wang JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                         2 LW
                                                                                          Matches
                                                                                                                                                                                                         RESULT 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                               AAB52657
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a monospecific antibody that binds specifically with an epitope defined by the amino acid sequence shown here. The peptide corresponds to a fragment of a trypsin-digested human fibrinogen and is designated fragment B(beta)123-127. The antibody can be used for detecting and measuring enzyme-mediated breakdown of fibrinogen and fibrin. The specific detection of cleavage products of fibrin and fibrinogen permits the accurate and precise determination of thrombotic activity in individuals. The products can be used to study thrombotic development and thrombolysis.
                                                                                                                                                 The present sequence represents an antigenic peptide of streptokinase. The specification describes a polypeptide which binds to a streptokinase-specific antibody and prevents the antibody binding to native streptokinase. The specification also describes a synthetic polypeptide (P1) comprising an epitope which binds to an streptokinase-specific antibody and reduces thrombolytic activity of streptokinase. P1 is used in thrombolytic therapy, and to prevent or treat glomerulonephritis and rheumatic fever.
                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New monospecific antibody to fibrin(ogen) - which can selectively bind matrix metalloproteinase-mediated cleavage fragments of fibrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antibody; epitope; trypsin; human; fibrinogen; detection; enzyme; cleavage; thrombosis; thrombus; thrombolysis.
                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                New polypeptides which bind streptokinase-specific antibodies useful in thrombolytic therapy
                                                                                                                                                                                                                                                                                                                                              65.2%; Score 15; DB 20; Length 5; 100.0%; Pred. No. 7.8e+05; Live 0; Mismatches 0; Indels
                                                                                                                   Example 1; Page 24; 44pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fibrinogen peptide B(beta)123-127.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW99381 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 34; 75pp; English
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Parhami-Seren B, Reed GL,
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                                  WPI; 1999-190113/16.
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Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 formyl peptide receptor family; FPR; inflammatory response up-regulation; chemoattractant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is a peptide fragment of T20/DP178. T20/DP178 is helical segment of the ectodomain of HIV-1 protein gp41. T20/DP178 is located in the carboxyl terminus of the gp41 ectodomain. gp41 plays a critical role in the fusion of HIV-1 and host cell membranes. T20/DP178 interacts with members of the formyl peptide receptor (FPR) family and thereby up-regulates an inflammatory response, and acts as a potent chemoattractant and activator of human peripheral blood phagocytes (but not T cells). The present peptide can be used to modulate an inflammatory response in a subject.
                                                                  Gaps
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partial-length T20/DP178 or T21/DP107 formyl peptide receptor
antagonist is used to modulate inflammation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gp41 ectodomain; HIV-1 fusion;
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65.2%; Score 15; DB 20; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus type 1.
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                                                                                                                                                                                                                                                                                                                        AAB52657 standard; Peptide; 5 AA.
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                                                                                                                                                February 12, 2003, 10:52:34; Search time 29 Seconds (without alignments) 4.405 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
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(c) 1993 - 2003 Compugen Ltd.
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US-09-201-396-7

US-10-105-930-70

US-10-105-930-70

US-09-903-412-51

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Maximum DB E
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GENERAL INFOGUEDATION

APPLICANT: Baxter, John D.

APPLICANT: Baxter, John D.

APPLICANT: Baxter, John D.

APPLICANT: Perg, Weijun

APPLICANT: Freq, Weijun

APPLICANT: Fletterick, Robert J.

APPLICANT: Wagner, Richard L.

APPLICANT: Wagner, Richard
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APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Partick
APPLICANT: Chene, Partick
APPLICANT: Chene, Partick
APPLICANT: Garcia-Eberoverria, Carlos
APPLICANT: Garcia-Eberoverria, Carlos
APPLICANT: Ruret, Pascal
TILE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPRENEUE: 4-2037/A/PCT
CURRENT APPLICATION NUMBER: U8/09/214,371B
CURRENT APPLICATION NUMBER: U8/09/214,371B
CURRENT APPLICATION NUMBER: U8/09/214,371B
FRIOR APPLICATION NUMBER: U8/09/214,371B
FRIOR FILING DATE: 1999-03-26
FRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
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                                      Sequence 74, P
Sequence 76, P
Sequence 78, P
Sequence 82, P
Sequence 84, P
Sequence 84, P
       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 82.6%; Score 19; DB 10; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 5; Conservative 0; Mismatches 0; Indels
US-09-903-412-69
US-09-903-412-74
US-09-903-412-76
US-09-903-412-78
US-09-903-412-82
US-09-903-412-82
US-09-903-412-84
US-09-903-412-84
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Patent No. US20020061539A1
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US-09-214-371-2
       SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                    Gaps
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APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE J. SCHLAUDER
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           D377/AP6D
                                                                                                                                                        LOCATION: (2)..(3)
OTHER INFORMATION: Xaa = Any Amino Acid
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; Patent No. US20020119447A1
; GENERAL INFORMATION:
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COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435435
ATTORNEY/AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PF
TELECOMMUNICATION INFORMATION:
TELEFHONE: 708-938-2655
TELEFAX: 708-938-2653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 708-938-2623 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 amino acids
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ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                   SEQ ID NO 3
LENGTH: 5
TYPE: PRT
ORGANISM: Homo sapiens
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GY: linear
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                                                                                                                                  NAME/KEY: SITE LOCATION: (2)...
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US-08-424-550B-228
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APPLICANT: Tomilinson, Ian M
APPLICANT: Winter, Gregory
TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands
FILE REFERENCE: 8039/10738
CURRENT APPLICATION NUMBER: US/09/968,561A
CURRENT FILING DATE: 2001-10-01
PRIOR FILING DATE: 1997-11-0-20
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: US 60/066,729
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: US 60/066,729
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: US 60/066,729
PRIOR FILING DATE: 1997-11-21
PRIOR APPLICATION NUMBER: US 60/066,739
PRIOR FILING DATE: 1998-10-20
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APPLICANT: Murphy, Andrew J.

APPLICANT: Murphy, Andrew J.

APPLICANT: Murphy, Andrew J.

APPLICANT: Paul, Joeremy

TITLE OF INVENTION: EFFECTORS

TITLE OF INVENTION: EFFECTORS

FILE REPERENCE: CPI-012CP9

CURRENT APPLICATION NUMBER: 08/09/201,396A

CURRENT APPLICATION NUMBER: 08/582,333

EARLIER APPLICATION NUMBER: 08/322,137

EARLIER PILING DATE: 1994-10-13

EARLIER PELING DATE: 1994-01-31

EARLIER PILING DATE: 1994-01-31

EARLIER PILING DATE: 1994-01-31

EARLIER PILING DATE: 1994-01-31

EARLIER APPLICATION NUMBER: 08/190,328

EARLIER PILING DATE: 1993-03-31

EARLIER APPLICATION NUMBER: 08/190,328

EARLIER PILING DATE: 1993-03-31

NUMBER OF SEQ ID NOS: 47

NUMBER OF SEQ ID NOS: 47

NUMBER OF SEQ ID NOS: 47
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50.0%; Pred. No. 1.2e+05;
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                              Sequence 217, Application US/09968561A patent No. US20020164642A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 350
SOFTWARE: PatentIn version 3.1
SEQ ID NO 217
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ORGANISM: Homo sapiens
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US-09-968-561A-217
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4 MW 5
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US-09-201-396-7
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GENERAL INVOKANTION:

APPLICANT: Tonlinson, lan M

APPLICANT: Finter, Gregory

TITLE OF INVERTION: Method to Screen Phage Display Libraries with Different Ligands

FILE REPERENCE: 8039/1073B

CURRENT PILING DATE: 2001-10-01

FRIOR PELING DATE: 1997-10-20

FRIOR APPLICATION NUMBER: US 60/065,248

FRIOR PELING DATE: 1997-11-3

FRIOR FILING DATE: 1997-11-21

FRIOR PELING DATE: 1997-11-21

FRIOR PELING DATE: 1998-10-20

FRIOR FILING DATE: 19
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Sequence 1.2.
Sequence 2.4.
Patent No. US20020061848al
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND TITLE OF INVENTION: UDAGNOSIS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121.469C8
CURRENT FILING DATE: 2010-04-23
NUMBER OF SEQ ID NOS: 599
SOUTHARE PASTSEQ FOR Windows Version 3.0/4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-841-132-9
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; ORGANISM: Homo sapiens
US-09-968-561A-211
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Best Local Similarity
Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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ઠે g 56.5%; Score 13; DB 9; Length 5;

Query Match

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Gaps
                             0; Gaps
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                                                                                                                                                                                                                                                                                                                       APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERBNCE: 0650-1-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: POT/JP00/06654
PRIOR APPLICATION NUMBER: UP 11-273358
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-17
NUMBER OF SEQ ID NOS: 77
SOFFWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        h Similarity 50.0%; Pred. No. 1.2e+05; 1; Conservative 1; Mismatches 0; Indels
ilarity 50.0%; Pred. No. 1.2e+05; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 51, Application US/09903412
; Publication No. US20030027319A1
; GENERAL INPORMATION:
; APPLICANT: Koide, Shohei
; TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
; FILE REFERENCE: 109.050US1
; CURRENT FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: US 60/217,474
; PRIOR FILING DATE: 2000-07-11
; PRIOR FILING DATE: 2000-07-11
; NUMBER OF SEQ ID NOS: 121
; SOFTWARE: FRASES for Windows Version 4.0
                                                                                                                                                                                                                         US-10-105-930-70
; Sequence 70, Application US/10105930
; Publication No. US20030009018A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 1; Conserva
Best Local Similarity
Matches 1; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Far
SEQ ID NO 51
                                                                                                      2 MW 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 ...
2 MW 3
                                                                           4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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APPLICANT: Winter, Greg
APPLICANT: Winter, Greg
APPLICANT: Winter, Greg
TITLE OF INVENTION: Methods for Selecting Functional Peptides
FILE REPRENCE: 3789/72916
CURRENT APPLICATION NUMBER: US/09/192,854
CURRENT FILING DATE: 1998-11-17
EARLIER PLING DATE: 1997-11-21
NUMBER OF SEQ ID NOS: 212
NUMBER OF SEQ ID NOS: 212
SOFTWARE: PatentIN Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 56.5%; Score 13; DB 10; Length 5; Best Local Similarity 50.0%; Pred. No. 1.2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                           APPLICANT: Koieda, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Bighth Street, Ste. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           109.034US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US/09/096,749A
                                           ; Sequence 51, Application US/09096749A; Patent No. US20020019517A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 120, Application US/09192854
; Patent No. US20020068276A1
                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INPORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.0
TELECOMMUNICATION INFORMATION:
TELEFROM (612) 373-6900
TELEFRAX: (612) 373-6900
TELEFRAX: (612) 373-6900
SEQUENCE CHARACTERISTICS: 51:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/C
FILING DATE: June 12, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                 Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FRAGMENT TYPE: i
; ORIGINAL SOURCE:
US-09-096-749A-51
                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                    55402
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RESULT 10
US-09-096-749A-51
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                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 LW 5
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Sequence 9, Application US/09924703

Patent No. US20020137898A1

GENERAL INFORMATION:

APPLICANT: FERUSSIA, BICE

APPLICANT: CLARK, STEVENC.

APPLICANT: CARRY, STEVENC.

APPLICANT: CLARK, STEVENC.

APPLICANT: MONG, CORDON G.

APPLICANT: MONG, CACRON G.

TITLE OF INVENTION: ANTIBODIES TO NATURAL KILLER STIMULATORY FACTOR

FILE REFERENCE: 01142.0142.0100

CURRENT APPLICANTON NUMBER: US/09/924,703

CURRENT FILING DATE: 1999-06-04

PRIOR FILING DATE: 1999-06-04

PRIOR FILING DATE: 1999-06-04

PRIOR FILING DATE: 1995-03-13

PRIOR FILING DATE: 1996-03-18

SECTUARES PATEANTIN VUMBER: 07/269,945

PRIOR FILING DATE: PATEANTIN VUMBER: 07/269,945
                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: peptide US-09-924-703-9
                                                                                                                                                   ;
0
                                                                                 Query Match 56.5%; Score 13; DB 10; Length 5; Best Local Similarity 50.0%; Pred. No. 1.2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 56.5%; Score 13; DB 10; Length 5; 50.0%; Pred. No. 1.2e+05; ative 1; Mismatches 0; Indels
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STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/08239765B
Patent No. US2002026650Al
GENERAL INFORMATION:
APPLICANT: Savakis, Charalambos
APPLICANT: Franz, Gerald H.
APPLICANT: Loukeris, Athanasios
TITLE OF INVENTION: Eukaryotic Transposable Element
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
; ORGANISM: Homo sapiens
US-09-192-854-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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US-08-239-765B-4
                                                                                                                                                                                                                                                                                                                                                                                US-09-924-703-9
                                                                                                                                                                                                       4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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US-09-096-749A-94

US-09-096-749A-94

Sequence 94, Application US/09096749A

Patent No. US20020019517A1

GENERAL INFORMATION:
APPLICANT: Koieda, Shohei

TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES

NUMBER OF SEQUENCES: 118

CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600

STREET: Minneapolis
STRIE: MN

COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 5;
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                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 09-MAY-1994
CLASSIFICATION : 435
PIGOR APPLICATION TABER: US/946,237
FILING DATE: 14-SEP-1992
ATTONNET/AGENT INFORMATION:
NAME: BLOOK, David B:
REGISTRATION NUMBER: 22,592
REFERENCE/POCKET NUMBER: 22,592
REGISTRATION NUMBER: 22,592
RESERVENCE/POCKET NUMBER: 22,592
REGISTRATION NUMBER: 22,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12; DB 8; I
Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: DISKETTE
COMPUTER: DISKETTE
COMPUTER: DISKETTE
COMPUTER: DISKETTE
COMPUTER: DISKETTE
APPLICATION NUMBER: US/09/096,749A
FILING DATE: JUNE 12, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEFAX: (612) 339-3061
                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 52.2%;
Best Local Similarity 50.0%;
Matches 1; Conservative
                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       , MOLECULE TYPE: peptide US-08-239-7658-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                               02173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 LW 5
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47.8%; Score 11; DB 8; Length 5; llarity 100.0%; Pred. No. 1.2e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                              SOFTWARE: Patentin Felease #1.0, Version #1.30
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUBER: US/08/484,409
FILING DATE: 07-JUN-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J. 31,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3: ABBOTT LABORATORIES D377/AP6D 100 ABBOTT PARK ROAD
                                                                                                                                                                                                                                                                                                                                                       690068.409CI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DANSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURBSH M. DESAI
APPLICANT: THOMAS P. LERRY
APPLICANT: ANTHONY SCOTT MUERHOFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 469, Application US/08424550B patent No. US20020119447A1 GENERAL INFORMATION:
                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 690
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JAMES C. ERKER
SHERI L. BUIJK
ISA K. MUSHAHWAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 amino acids
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                                                                                 COMPUTER READABLE FORM:
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                       Washington
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nes 1; Conserv
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                                                               98104-7092
Seattle
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APPLICANT:
APPLICANT:
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APPLICANT: PTITSTN, Leonid R
APPLICANT: PTITSTN, Leonid R
APPLICANT: PTITSTN, Leonid R
APPLICANT: SMIRNOV, Sergey V
APPLICANT: SMIRNOV, Yalia G
APPLICANT: YAMPOLSKAYA, Tatyana A
APPLICANT: TEONOVA, Tatyana V
APPLICANT: GUSYATINER, Mikhail M
TITLE OF INVENTION: NEW MUTANT-ACETYLGLUTAMATE SYNTHASE AND METHOD FOR L-ARGININE PRC
FILE REFERENCE: 209873USC
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: USSIAN FEDERATION 2000116481
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2001-06-15
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PATENTIN VERSION 3.1
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Patent No. US20020076412A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Steining, Lewrence
APPLICANT: Steining, Lowernce
APPLICANT: Zamvil, Scott
ITILE OF INVENTION: METHODS FOR MODULATING THE IMMUNE SYSTEM
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           52.2%; Score 12; DB 10; Length 5; 50.0%; Pred. No. 1.2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                   52.2%; Score 12; DB 10; Length 5; 50.0%; Pred. No. 1.2e+05; tive 1; Mismatches 0; Indels
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    94:
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Best Local Similarity 50.09
Matches 1; Conservative
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                     SEQUENCE CHARACTERISTICS:
  INFORMATION FOR SEQ ID NO:
                                       LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                           Sirving
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                    single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
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LENGTH: 5
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; NAME/KEY: Site
; LOCATION: (3)
: OTHER INFERMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
US-09-764-884-2
                                                                                                            47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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; Patent No. US20020160949A1
                                                                                                                                                                                                                                                                                                           ; Sequence 15, Application US/09895593 ; Patent No. US20020160949A1
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                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Pandey, Akhilesh
APPLICANT: Ozaki, Katsutoshi
                                                                                                                               Best Local Similarity 100. Matches 1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14, Application US/09995749A

Patent No. US20020155568A1

GENERAL INFORMATION:
APPLICANT: VAN GEBL-SCHUTTEN, GERRITDINA HENDRIKA
APPLICANT: DIJKHUIZEN, LUBBERT
APPLICANT: LEER, ROBERT-JAN
TITLE OF INVENTION: NOVEL GLUCOSYLTRANSFERASES
FILE REFERENCE: BO43188-CIP
CURRENT APLICATION NUMBER: US/09/995,749A
CURRENT FILING DATE: 2001-11-29
PRIOR APPLICATION NUMBER: 09/604,957
PRIOR APPLICATION NUMBER: EPO 00201871.1
PRIOR APLICATION NUMBER: EPO 00201871.1
PRIOR PILING DATE: 2000-06-28
PRIOR APPLICATION NUMBER: EPO 00201871.1
PRIOR PILING DATE: 2000-06-28

NUMBER OF SEQ ID NOS: 19
SOFTWARE: PatentIN Ver. 2.1
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Neisseria polysaccharea
US-09-995-749A-14
                                                                  TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 469:
SEQUENCE CHARACTER.STICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                        1; Conservative
                                          TELEPHONE: 708-22
TELEPHONE: 708-938-2623
                                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: protein
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ORGANISM: Homo sapiens
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XGY: linear
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Best Local Similarity
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US-09-995-749A-14
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LENGTH: 5
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APPLICANT: Levin, Resconting APPLICANT: Levin, Recent D.
APPLICANT: Levin, Steven D.
APPLICANT: Levin, Addrew G.
APPLICANT: Ziegler, Steven F.
APPLICANT: Ziegler, Steven F.
APPLICANT: Leonard, Warren J.
APPLICANT: Lodish, Harvey F.
TITLE OF INVENTION: Thymic Stromal Lymphopoietin Receptor Molecules and TITLE OF INVENTION: Uses Thereof FILE REFERENCE: 00-514-E
CURRENT APPLICATION NUMBER: US/09/895,593
CURRENT FILING DATE: 2001-06-28
PRIOR APPLICATION NUMBER: 60/215,658
PRIOR PLING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
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GENERAL INFORMATION:
APPLICANT: Pandey, Akhilesh
APPLICANT: Ozafi, Katsutoshi
APPLICANT: Baumann, Heinz
APPLICANT: Levin, Steven D.
APPLICANT: Leonar, Steven F.
APPLICANT: Leonard, Warren J.
APPLICANT: Leonard, Warren J.
TITLE OF INVENTION: Thymic Stromal Lymphopoietin Receptor Molecules and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps

; LOCATION: (3)
; OTHER INFORMATION: "Xaa" can be any naturally occurring amino acid
US-09-895-593-15

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                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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APPLICANT: TSUNEMARI, TOSHIAKI
APPLICANT: TSUNEMARI, TOSHIAKI
TITLE OF INVENTION: CACHEXIA REMEDY
FILE REPERBROKE: 04833-0038
CURRENT APPLICATION NUMBER: US/09/423,800
CURRENT FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: PCT/JP98/02116
PRIOR PLILING DATE: 1998-05-13
PRIOR PLILING DATE: 1998-05-15
PRIOR PLILING DATE: 1997-07-18
PRIOR PLILING DATE: 1997-07-18
             CURRENT APPLICATION NUMBER: US/09/895,593
CURRENT FILING DATE: 2001-06-28
PRIOR APPLICATION NUMBER: 60/215,658
PRIOR FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.0
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; Patent No. US20020165363A1
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FILE REFERENCE: 00-514-E
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US-09-423-800-62
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APPLICANT: SATO, K
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US-09-764-857-2
                                                                                                                                                                      SEQ ID NO 16
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LOCATION: (3) OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
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Patent No. US20020165137A1;
GENERAL INFORMATION:
APPLICANT: RUben et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies;
FILE REFERENCE: PA127P1
CURRENT APPLICATION NUMBER: US/09/860,670
CURRENT PILING DATE: 2001-05-21;
Prior application data removed - consult PALM or file wrapper;
NUMBER OF SEQ ID NOS: 289
SOSTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                    47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: ROSEn et al.
TITLE OF INVENSION: 70 Human Secreted Proteins FILE REFERENCE: PZ001P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: US/09/981,876
PRIOR APPLICATION NUMBER: 09/148,545
PRIOR PELING DATE: 1998-09-04
PRIOR PELING DATE: 1998-09-04
PRIOR PELING DATE: 1997-03-07
PRIOR PELING DATE: 1997-03-07
PRIOR PILING DATE: 1997-03-07
PRIOR PILING DATE: 1997-03-07
PRIOR PILING DATE: 1997-03-07
PRIOR PELING DATE: 1997-03-07
PRIOR APPLICATION NUMBER: 60/040,161
PRIOR APPLICATION NUMBER: 60/040,626
PRIOR APPLICATION NUMBER: 60/040,626
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; Patent No. US20020164669A1
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 5
                                                                                                                                                                                                                                                                                                     Conservative
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ORGANISM: Homo sapiens
                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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nes 1; Conserva
                                                                                                                                    NAME/KEY: Site
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LOCATION: (3)
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US-09-981-876-2
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LENGTH: 5
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R FILING DATE: 1997-03-07
R APPLICATION NUMBER: 60/040,163
R FILING DATE: 1997-03-07
R R APPLICATION NUMBER: 60/047,615
R FILING DATE: 1997-05-23
R RAPPLICATION NUMBER: 60/047,600 FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,334
FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,336 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,587 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,597 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,633 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,583 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,617 APPLICATION NUMBER: 60/047,618 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,503 APPLICATION NUMBER: 60/047,592 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,584 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,492 APPLICATION NUMBER: 60/047,598 APPLICATION NUMBER: 60/047,582 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,596 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,612 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,632 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,580 APPLICATION NUMBER: 60/043,568 APPLICATION NUMBER: 60/043,314 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,569 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,311 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,313 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,672 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/047,502 APPLICATION NUMBER: 60/047,581 APPLICATION NUMBER: 60/047,500 APPLICATION NUMBER: 60/047,613 APPLICATION NUMBER: 60/047, 601 APPLICATION NUMBER: 60/043,671 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,674 APPLICATION NUMBER: 60/043,312 ILING DATE: 1997-05-23 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 ILING DATE: 1997-05-23 LING DATE: 1997-04-11 ILING DATE: 1997-04-11 1997-04-11 ILING DATE: 1997-04-1 ILING DATE: 1997-04-1 ILING DATE: RRILIORA RELIGIORA RELIGIO

FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,911
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,636 APPLICATION NUMBER: 60/056,892 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/047,595 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/057,761 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/048,974 FILING DATE: 1997-06-06 APPLICATION NUMBER: 60/056,886 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,877 FILING DATE: 1997-08-22
APPLICATION WUMBER: 60/056,889
FILING DATE: 1997-08-23
APPLICATION NUMBER: 60/056,893 APPLICATION NUMBER: 60/056,888 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,879 APPLICATION NUMBER: 60/043,315 APPLICATION NUMBER: 60/056,630 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,878 APPLICATION NUMBER: 60/056,662 FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,882 APPLICATION NUMBER: 60/056,637 FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,874 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,910 APPLICATION NUMBER: 60/047,585 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,614 APPLICATION NUMBER: 60/056,872 APPLICATION NUMBER: 60/056,903 APPLICATION NUMBER: 60/056,880 APPLICATION NUMBER: 60/056,894 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,864 APPLICATION NUMBER: 60/056,631 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,845 FILING DATE: 05-Sep-1997 APPLICATION NUMBER: 60/047,599 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,588 APPLICATION NUMBER: 60/047,586 APPLICATION NUMBER: 60/047,590 APPLICATION NUMBER: 60/047,594 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,589 APPLICATION NUMBER: 60/047,593 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,578 FILING DATE: 1997-08-22 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 1997-05-23 FILING DATE: 1997-05-23 1997-05-23 PRILICA REPUBLICA REPUBLIC

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TITLE OF INVENTION: 87 Human Secreted Proteins
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CURRENT APPLICATION NUMBER: US/09/984,245
CURRENT FILING DATE: 2001-10-29
PRIOR PPLICATION NUMBER: US/09/984,245
PRIOR FILING DATE: 1998-09-17
PRIOR APPLICATION NUMBER: OF/US98/05311
PRIOR PILING DATE: 1998-03-19
PRIOR PILING DATE: 1998-03-19
PRIOR PILING DATE: 1997-03-21
PRIOR PILING DATE: 1997-03-21
PRIOR PILING DATE: 1997-03-21
PRIOR PLING DATE: 1997-03-30
PRIOR PLING DATE: 1997-05-30
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PRIOR PLILING DATE: 1997-05-23
PRIOR PLILING DATE: 1997-04-11
PRIOR PLILING DATE: 1997-04-11
PRIOR FILING DATE: 1997-04-12
PRIOR FILING DATE: 1997-08-22
PRIOR PELING DATE: 1997-08-22
PRIOR PLILING DATE: 1997-08-22
PRIOR PELING DATE: 1997-08-22
PRIOR PELING DATE: 1997-08-22
PRIOR PLILING DATE: 1997-08-22
PRIOR PRILING DATE: 1997-08-22
PRIOR PLILING DATE: 1997-08-22
FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/043,576
                                                                                  FILING DATE: 1997-04-11
APPLICATION NUMBER: 60/047,501
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Sequence 2, Application US/09984245
Patent No. US20020165374A1
GENERAL INFORMATION:
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OTHER INFORMATION: Xaa equals any of the L-amino acids commonly found in naturally OTHER INFORMATION: proteins
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PT232
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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PRIOR APPLICATION NUMBER: US 60/048,160
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR PLING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/048,154
PRIOR FILING DATE: 1997-08-09
PRIOR FILING DATE: 1997-08-09
PRIOR FILING DATE: 1997-08-19
PRIOR FILING DATE: 1997-08-19
PRIOR FILING DATE: 1997-08-19
PRIOR PLING DATE: 1997-08-19
PRIOR FILING DATE: 1997-08-19
PRIOR FILING DATE: 1997-08-19
PRIOR PLING DATE: 1997-08-19
PRIOR PLING DATE: 1997-08-19
PRIOR PLING DATE: 1997-08-19
PRIOR PLING DATE: 1997-10-02
NUMBER OF SEQ ID NOS: 343
SEQ ID NO 2
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,095
                           PILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/050,937
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,069
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/048,355
APPLICATION NUMBER: US 60/048,135
                                                                                                              APPLICATION NUMBER: US 60/048,187
                                                                                                                                                                                                FILING DATE: 1997-05-30 APPLICATION NUMBER: US 60/048,352
                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/048,186
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; Patent No. US20020168711A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 1997-05-30
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Matches 1; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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LENGTH: 5
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NAME/KEY: Site
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US-09-764-904-2
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Fatent No. US20020168684A1
GENERAL INFORMATION:
APPLICANT: Cell Signaling Technology, Inc.
APPLICANT: COMB, Michael J.
APPLICANT: COMB, Michael J.
APPLICANT: COMB, Michael J.
APPLICANT: CHI Signaling Technology, Inc.
APPLICANT: COMB, Michael J.
APPLICANT: COMB, Michael J.
APPLICANT: TAN, Yi
TITLE OF INVENTION: PRODUCTION OF MOTIF-SPECIFIC AND CONTEXT-INDEPENDENT ANTIBODIES USTILE OF INVENTION: PREPRIED LIBRARIES AS ANTIGENS
FILE REFERENCE: CST-138 CIP2
CURRENT APPLICATION NUMBER: US/10/014,485A
CURRENT APPLICATION NUMBER: US 09/148,712
PRIOR FILING DATE: 1998-09-04
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 145
SOFFWARE PATENTIN VERSION 3.1
                    LOCATION: (3)
CTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids US-09-764-868-2
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: LOCATION: (3) .. (3)

: COTER INCRMATION: PHOSPHORYLATION; threonine at position 3 is phosphorylated US-10-014-485A-79
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Sequence 42, Application US/10104019

Batent No. US20020168379A1

GENERAL INFORMATION:
APPLICANT: MENG, Xiang-Jin
APPLICANT: MENG, Xiang-Jin
APPLICANT: MENG Xiang-Jin
APPLICANT: MENG Xiang-Jin
APPLICANT: MENG Xiang-Jin
APPLICANT: Patrick
TITLE OF INVENTION: PROTEINS ENCODED BY POLYNUCLEIC ACIDS OF PORCINE
TITLE OF INVENTION: PROTEINS ENCODED BY POLYNUCLEIC ACIDS OF PORCINE
FILE REFERENCE: 4625-0039-55X CIP
CURRENT APPLICATION NUMBER: US/10/104,019
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                                                                                                                                                                     0; Indels
                                                                                                                     Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 1998-02-06
PRIOR APPLICATION NUMBER: 08/301,435
PRIOR FILING DATE: 1994-09-01
PRIOR PLING DATE: 1994-09-01
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NAME/KEY: Site
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US-10-014-485A-79
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US-10-104-019-42
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; LOCATION: (3)
; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
US-09-764-904-2
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Fatent No. US20020173454A1

GENERAL INFORMATION:

APPLICANT: Rosen et al.

TITLE OP INVENTION: Nucleic Acids, Proteins, and Antibodies

FILE REFERENCE: PA122

CURRENT FILING DATE: 2001-01-17

Prior application data removed - consult PALM or file wrapper

NUMBER OF SEQ ID NOS: 137

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 2

LENGTH: 5
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                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: Peptide US-10-104-019-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                 Query Match

47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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Sequence 2, Application US/09852797
Patent No. US20020172994A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: 28 Human Secreted Proteins
FILE REFERENCE: PZ003P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/852,797
CURRENT FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 60/265,583
PRIOR FILING DATE: 2001-02-02
PRIOR PILING DATE: 1998-09-11
PRIOR PILING DATE: 1998-03-12
PRIOR PELICATION NUMBER: PCT/US98/04858
PRIOR PELING DATE: 1998-03-12
PRIOR PELING DATE: 1998-03-14
PRIOR PELING DATE: 1997-03-14
PRIOR PELING DATE: 1997-03-14
PRIOR FILING DATE: 1993-10-05
PRIOR APPLICATION NUMBER: 07/969,071
PRIOR FILING DATE: 1992-10-30
NUMBER OF SEQ ID NOS: 108
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 42
SEROID NO 42
                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Homo sapiens
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Sequence 8, Application US/10035349
; Sequence 8, Application US/10035349
; Patent No. US20020172961A1
; GENERAL INFORMATION:
    APPLICANT: Schneider, Luke V.
    APPLICANT: Hall, Michael P.
    APPLICANT: Hall, Michael P.
    APPLICANT: Target Discovery, Inc.
    TITLE OF INVENTION: Sequences
    FILE REFERENCE: 020444-00071008
; CURRENT APPLICATION NUMBER: US/10/035,349
; CURRENT PILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: US 60/242,165
; PRIOR APPLICATION NUMBER: US 60/242,165
; PRIOR APPLICATION NUMBER: US 60/242,398
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                           APPLICANT: MILY TARGET, ROBERT APPLICANT: Petesch, Robert APPLICANT: Petesch, Robert APPLICANT: Target Discovery, Inc.
TITLE OF INVENTION: Mess Defect Labeling for the Determination of Oligomer TITLE OF INVENTION: Sequence6
FILE REPRENDE: 02044-000710US
CURRENT APPLICATION NUMBER: US 60/242,165
PRIOR APPLICATION NUMBER: US 60/242,165
PRIOR APPLICATION NUMBER: US 60/242,398
PRIOR RILING DATE: 2000-10-19
PRIOR FILING DATE: 2000-10-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence:competing 5 OTHER INFORMATION: residue false sequence, top ranked sequence at OTHER INFORMATION: five residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Description of Artificial Sequence:competing 5 OTHER INFORMATION: residue false sequence, top ranked sequence at OTHER INFORMATION: five residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Xaa = Gly labeled with 5-bromo-3-pyridylacetic OTHER INFORMATION: acid (5-Br-3-PAA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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                                                    ; Sequence 2, Application US/10035349; Patent No. US20020172961A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                          APPLICANT: Schneider, Luke V. APPLICANT: Hall, Michael P.
                                                                                                                 GENERAL INFORMATION:
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NAME/KEY: MOD RES
                                 US-10-035-349-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids US-09-852-797-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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APPLICANT: Campos, Reinaldo
APPLICANT: Campos, Reinaldo
APPLICANT: No. US200201013633Alogaki, Hiroyuki
APPLICANT: No. US200201013633Alogaki, Hiroyuki
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of Phenylalanine Ammonia-Lyase (PAL)
TITLE OF INVENTION: Characterization of Phenylalanine Ammonia-Lyase (PAL)
FILE REPERBUCE: 023070-1245000 Lettuce Tissue
CURRENT APPLICATION NUMBER: US/09/964,992A
CURRENT FILING DATE: 2000-09-26
PRIOR PRILING DATE: 2000-09-26
PRIOR FILING DATE: 2000-09-26
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                              PRIOR FILING DATE: 1997-05-30

PRIOR APPLICATION NUMBER: 60/048,100

PRIOR FILING DATE: 1997-05-30

PRIOR PILING DATE: 1997-05-30

PRIOR PILING DATE: 1997-09-05

PRIOR APPLICATION NUMBER: 60/048,189

PRIOR PILING DATE: 1997-09-05

PRIOR PILING DATE: 1997-09-05

PRIOR FILING DATE: 1997-09-05

PRIOR FILING DATE: 1997-05-06

PRIOR FILING DATE: 1997-12-19

NUMBER OF SEQ ID NOS: 118

SEQ ID NOS: 118

SEQ ID NOS

LENGTH: 5
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                                 PRIOR APPLICATION NUMBER: 60/050,934
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ORGANISM: Artificial Sequence
PRIOR FILING DATE: 1997-03-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 1; Conserva
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US-10-035-349-8

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TITLE OF INVENTION: Apoptotic Protease Mch6, Nucleic Acids
Encoding Same and Methods of Use
                                                                                                                                                                                                     APPLICANT: Alnemri, Emad S.
Pernandes-Alnemri, Teresa
Litwack, Gerald
TITLE OF INVENTION: Applotic Protease Mch6, Nucleic Acids
Encoding Same and Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: United States
ZIP: 92122
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: DEAD PC DOWNSTED: COMPATING SYSTEM: PC-DOS/MS-DOS
COFRANTING SYSTEM: PC-DOS/MS-DOS
COFRANTS: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-Jan-2002
CLASSIFICATION: Unknown>
PRIOR APPLICATION NUMBER: US/08/865,579
FILING DATE: 29-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31, 815
REFERENCE/DOCKET NUMBER: P-ID 2180
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 63, Application US/10059749; Publication No. US20020183504A1; GENERAL INFORMATION: APPLICANT: Alnemri, Emad S.; Fernandes-Alnemri, Teresa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 57:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-9849
                                                                                                                            ; Sequence 57, Application US/10059749; Publication No. US20020183504A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 87
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
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                                                                                            RESULT 37
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US-09-828-272A-6
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APPLICANT: CATANIA, Anna P
TITLS OF INVENTION: USE OF KPV TRIPEPTIDE FOR DERMATOLOGICAL DISORDERS
FILE REFERENCE: 259/058
CURRENT PAPLICATION UNDERR: US/09/828,272A
CURRENT FILING DATE: 2001-04-06
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.1
SEQ ID NO 6
LENGTH: 5
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                                           0; Indels
47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; Live 0; Mismatches 0; Indels
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US-10-042-141-2
Sequence 2, Application US/10042141
Sequence 2, Application No. US20020183503A1
GENERAL INFORMATION:
TITLE OF INVENTION: 26 Human secreted proteins
FILE REFERENCE: PZ040P1
CURRENT PELLON NUMBER: US/10/042,141
CURRENT FILING DATE: 2002-01-11
PRIOR APPLICATION NUMBER: 09/726,643
FRIOR APPLICATION NUMBER: PCT/US00/15187
PRIOR FILING DATE: 2000-06-02
PRIOR FILING DATE: 2000-06-02
PRIOR FILING DATE: 1999-06-07
NUMBER OF SEQ ID NOS: 190
                                                                                                                                                                                                                                                        ; Sequence 6, Application US/09828272A; Publication No. US20020183255A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
  Query Match
Best Local Similarity 100.0
Matches 1; Conservative
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                                                                                                                                                                                                            RESULT 35
US-09-828-272A-6
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Publication No. US20020183504A1
GENERAL INFORMATION:
APPLICANT: Alnemri, Emad S.
Fernandes-Alnemri, Teresa
Litwack, Gerald
TITLE OF INVENTION: Apoptotic Protease Mch6, Nucleic Acids
Encoding Same and Methods of Use
                                                                                         ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-Jan_2002
CLASSIFICATION : vURNOWN>
PRIOR APPLICATION DATA:
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSE: Campbell & Flores LLP
STREET: 4770 La Jolla Village Drive, Suite 700
CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AGENTE: 29-MAY-1997
ATTORNEY AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
FERERENCE/COCKET NUMBER: P-ID 2180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/865,579
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FILING DATE: 29-MAY-1997
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-0an-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear;
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 63:
US-10-059-749-63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (619) 535-9849
                                                                      COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: California
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 63:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 87
                                                 STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                            CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-059-749-69
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APPLICANT: Tepper, Robert I.
APPLICANT: Tepper, Robert I.
APPLICANT: Culpepper, Janice A.
APPLICANT: Culpepper, Janice A.
APPLICANT: White, David W.
TITLE OF INVENTION: THE OB RECEPTOR AND METHODS FOR
TITLE OF INVENTION: THE DIAGNOSIS AND TREATMENT OF BODY WEIGHT DISORDERS,
TITLE OF INVENTION: INCLUDING OBESITY AND CACHEXIA
NUMBER OF SEQUENCES: 50
CORRESSORE Fish & Richardson, P.C.
STREET: 225 Franklin Street
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                                                                                                                                                                                                                                                                                                                                                                                      Length 5;
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47.8%; Score 11; DB 9; Le
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0;
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: RASTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENY APPLICATION DATA:
FILING DATE: 2002-FEB-19
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-ID 2180
TELECOMMUNICATION INFORMATION:
TELEPRAK: (619) 535-9001
TELEFRAK: (619) 535-9849
                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 69:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/84,564
FILING DATE: 28-MAY-1997
APPLICATION NUMBER: 08/708,123
FILING DATE: 03-SEP-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 22-ARR-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 22-DAY-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 28-DEC-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 18-DEC-1996
APPLICATION NUMBER: 08/59,455
FILING DATE: 11-DEC-1995
APPLICATION NUMBER: 08/59,455
FILING DATE: 08-DEC-1995
APPLICATION NUMBER: 08/59,455
FILING DATE: 08-DEC-1995
APPLICATION NUMBER: 08-56,652
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APPLICATION NUMBER: 08/562,663
FILING DATE: 27-NOV-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 6, Application US/10079625; Publication No. US20020182676A1; GENERAL INFORMATION:
                                                                                                                                                                                                         LENGTH: 5 amino acids
                                                                                                                                                     INFORMATION FOR SEQ ID NO: 69: SEQUENCE CHARACTERISTICS:
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US-10-079-625-6
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NAME: Meiklejohn, Ph.D., Anita L.

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APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
TITLE OF INVENTION: Component of Stem Bromelain (as amended)
TITLE OF INVENTION: Component of Stem Bromelain (as amended)
FILE REPERBENCE: 0623.0760002
CURRENT APPLICATION NUMBER: US/10/029,301
CURRENT FILING DATE: 1999-08-25
FRIOR PILING DATE: 1999-08-25
FRIOR APPLICATION NUMBER: CB 9704252.7
FRIOR APPLICATION NUMBER: GB 970452.7
FRIOR PILING DATE: 1997-02-28
FRIOR FILING DATE: 1997-02-25
FRIOR FILING DATE: 1997-02-25
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                              Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: METHODS OF TREATING BLADDER DISORDERS
FILE REFERENCE: 08191-022001
CURRENT APPLICATION NUMBER: US/10/074,956
CURRENT FILING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: 60/268,175
                          07334/019002
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Publication No. US20020188107A1
GENERAL INFORMATION:
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Publication No. US20020193332A1
GENERAL INFORMATION:
                  REFERENCE/DOCKET NUMBER: 07334
TELECOMMUNICATION INPORMATION:
TELEPHONE: 617-542-8906
TELERAX: 200154
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
REGISTRATION NUMBER: 35,283
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SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-10-079-625-6
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                                                                                                                                                                                                                                                             unknown
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TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: RUDGE et al.

TITLE OF INVENTION: Cytokine Receptor Common Gamma Chain Like
FILE REFERENCE: PF466P2
CURRENT APPLICATION WUBBER: US/10/078,059
CURRENT FILING DATE: 2002-02-20
FRIOR APPLICATION WUBBER: 60/269,876
PRIOR PLING DATE: 2000-03-21
PRIOR PLING DATE: 2000-08-17
PRIOR APPLICATION WUBBER: PCT/US00/22493
PRIOR PLING DATE: 2000-08-17
PRIOR APPLICATION WUBBER: 09/376,430
PRIOR PLING DATE: 1999-08-18
PRIOR PLING DATE: 1999-03-05
PRIOR PLING DATE: 1999-03-05
PRIOR PLING DATE: 1998-03-05
PRIOR APPLICATION WUBBER: 60/086,505
PRIOR APPLICATION WUBBER: 60/078,563
PRIOR PLING DATE: 1998-03-19
WUBBER OF SEQ 1D NOS: 32
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                                                                                                                                                                                                                                 Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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APPLICANT: Raesen, Guido, Christiaan
TITLE OF INVENTION: Treatment of Allergic Rhinitis
FILE REFERENCE: 2468-1-004
CURRENT APPLICATION NUMBER: US/10/097,195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (3)
OTHER INFORMATION: Xaa equals any amino acid
                      NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-078-059-5
; Sequence 5, Application US/10078059
; Publication No. US20020193305Al
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 3, Application US/10087195; Publication No. US20020193306Al; GENERAL INFORMATION:
PRIOR FILING DATE: 2001-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: SITE
                                                                                                                                                                                  US-10-074-956-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-10-078-059-5
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TITLE OF INVENTION: 90 Human Secreted Proteins
FILE REPERENCE: P2013P1
CURRENT APPLICATION NUMBER: US/09/774,639
CURRENT FILING DATE: 2001-07-09
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/244,112
PRIOR FILING DATE: EARLIER FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 371
                                                                                                                                                                                                                                                                                                                            Query Match

47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match

47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 1999-05-27
EARLIER APPLICATION NUMBER: 60/099,805
EARLIER APPLICATION NUMBER: 60/099,805
EARLIER FILING DATE: 1998-09-10
EARLIER PILING DATE: 1999-04-30
NUMBER OF SEQ ID NOS: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: x equals any amino acid US-09-320-713-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Reinhard Ebner
APPLICANT: Steven M. Ruben
TITLE OF INVENTION: INTERLEUKINS-21 AND 22
FILE REFERENCE: PF470
CURRENT FILING DATE: 2002-03-01
PRICA APPLICATION NUMBER: PCT/GB00/03287
PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: 9920673.2
PRIOR FILING DATE: 1999-09-01
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 14, Application US/09320713
Publication No. US20030003545A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                           TYPE: PRT ORGANISM: arthropod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: SITE
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US-09-774-639-2
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LENGTH: 5
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TITLE OF INVENTION: 31 Human secreted proteins
TITLE OF INVENTION: 31 Human secreted proteins
FILE REFERENCE: P2026P1
CURRENT PILING DATE: 2001-03-06
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/393,022
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/077,714
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/077,714
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: EARLIER FILING DATE: 1998-03-12
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PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
NUMBER OF SEQ ID NOS: 185
SOFTWARE: PATENTING DATE: 2.0
SEQ ID NO 2
LENGTH: 5
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Publication No. US20030009018A1
GENERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105U31
CURRENT APPLICATION NUMBER: US/10/105,930
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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SOFTWARE: Patentin Ver. 2.0
                                                                            TYPE: PRT
ORGANISM: Homo sapiens
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NAME/KEY: Site
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                       SEQ ID NO 2
LENGTH: 5
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US-10-105-930-59

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squence 59, Application US/10105930

special No. US20030009018A1

squence 59, Application No. US20030009018A1iko

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT FILING DATE: 2002-03-25

PRIOR APPLICATION NUMBER: PCT/JP00/06654

PRIOR APPLICATION NUMBER: JP 2000-240397

PRIOR FILING DATE: 2000-08-03

PRIOR FILING DATE: 1999-09-27

NUMBER OF SGD ID NOS: 77

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 5
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Publication No. US20030009018A1
GENERAL INFORMATION:
APPLICANT: Maeda, Masteugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFRENCE: 06501-105U31
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: DT/JP00/06654
PRIOR APPLICATION NUMBER: JP02-03-27
PRIOR APPLICATION NUMBER: JP02-03-27
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 9; Le
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0;
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: DP 2000-240397
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-03
PRIOR FILING DATE: 199-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-105-930-58
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US-10-105-930-60
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APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Mouse, US20030099018Aliko
TITLE OP INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-1054S1
CURRENT PILING DATE: 2002-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: DCT/JP00/06654
PRIOR APPLICATION NUMBER: DP 2000-240397
PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR PILING DATE: 2000-08-03
PRIOR FILING DATE: 1999-09-77
NUMBER OF SEQ 1D NOS: 77
SOFTWARE PAECES FOR WINDOWS Version 4.0
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Publication No. US2003009018A1
GENERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOPTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; LOCATION: 3; OTHER INFORMATION: Xaa = any amino acid US-10-105-930-21
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                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Exemplary motif
        CURRENT FILING DATE: 2002-03-25
                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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; ORGANISM: Homo sapiens
US-10-105-930-57
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LENGTH: 5
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; SEQ ID NO 62
; LENGTH: 5
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-105-930-62
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LENGTH: 5
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APPLICANT: Masatsugu

APPLICANT: Yaguchi, No. US20030009018Aliko

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

FILE REFERENCE: 06501-105US1

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/JP00/06654

PRIOR APPLICATION NUMBER: PCT/JP00/06654

PRIOR PLING DATE: 2000-09-27

PRIOR PLING DATE: 1000-08-03

PRIOR PLING DATE: 1999-09-27

NUMBER OF SOOD NOS: 77

NUMBER OF SOOD NOS: 77

SOFTWARE PASSEQ for Windows Version 4.0
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; Publication No. US2003009018A1
; GENERAL INFORMATION:
APPLICANT: Masatsugu
APPLICANT: Yaquchi, No. US2003009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 65601-105481
; CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT APPLICATION NUMBER: D002-03-25
; PRIOR APPLICATION NUMBER: PCT/JP00/06654
; PRIOR APPLICATION NUMBER: D11-273358
; PRIOR APPLICATION NUMBER: JP 11-273358
; PRIOR APPLICATION NUMBER: JP 11-273358
; PRIOR APPLICATION NUMBER: JP 11-273358
; RIOR APPLICATION NUMBER: JP 11-273358
; NUMBER OF SEQ ID NOS: 77
; SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 61, Application US/10105930; Publication No. US20030009018A1; GENERAL INFORMATION:
                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
US-10-105-930-60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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US-10-105-390-68
Sequence 68, Application US/10105930
Sequence 68, Application US/10105930
Fublication No. US20030009018A1
Sequence 10 No. US20030009018A1
SERENAL INFORMATION:
APPLICANT: Maeda, Masatsugu
TITLE OF INVENTION: NOVEL HEMOPOLETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105U31
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT PLING DATE: 2002-03-25
FRIOR RELIGNED DATE: 2000-09-27
FRIOR PELING DATE: 2000-09-27
FRIOR FILING DATE: 2000-09-03
FRIOR FILING DATE: 2000-09-03
FRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NOS: 77
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APPLICANT: Waeda, Masateugu

APPLICANT: Yaguchi, No. US20030009018Aliko

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

FILE REPERENCE: 06501-105US1

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT APPLICATION NUMBER: PCT/JP00/06654

PRIOR PILING DATE: 2000-09-27

PRIOR PELLING DATE: 2000-09-27

PRIOR PILING DATE: 1999-09-27

PRIOR PILING DATE: 1999-09-27

PRIOR APPLICATION NUMBER: JP 11-273358

PRIOR APPLICATION NUMBER: JP 11-273358

NUMBER OF SEQ ID NOS: 77

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                      US-10-105-930-67
; Sequence 67, Application US/10105930
; Publication No. US20030009018A1
; GENERAL INFORMATION:
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US-10-105-930-68
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CORGANISM: Homo sapiens
US-10-105-930-67
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US-10-105-930-66

Sequence 66, Application US/10105930

Publication No. US2003009018A1

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 06501-105US1

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT PILING DATE: 2002-03-25

PRIOR APPLICATION NUMBER: PCT/JP00/06654

PRIOR APPLICATION NUMBER: UP 2000-240397

PRIOR APPLICATION NUMBER: UP 11-273358

PRIOR FILING DATE: 2000-09-27

PRIOR FILING DATE: 1999-09-27

NUMBER OF SEQ ID NOS: 77

SOFTWARE: FASTERED FOR WINDOW VERSION 4.0

SEQ ID NOS: 77
                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Masateugu
APPLICANT: Masateugu
APPLICANT: Masateugu
APPLICANT: Masateugu
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 66501-10543
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT APPLICATION NUMBER: PGT/JP00/06654
PRIOR APPLICATION NUMBER: PGT/JP00/06654
PRIOR APPLICATION NUMBER: J0 2000-240397
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-09
PRIOR PILING DATE: 1999-09-09
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                   0; Indels
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                   47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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; ORGANISM: Homo sapiens
US-10-105-930-66
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US-10-105-930-65
                   Query Match
Best Local Similarity
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                                                                     Matches
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APPLICANT: Maeda, Masatsugu

APPLICANT: Maeda, Masatsugu

APPLICANT: Maeda, Mosatsugu

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

FILE REFERENCE: 0650.1-05081

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT FILING DATE: 2002-03-25

PRIOR APPLICATION NUMBER: PCT/JP00/06654

PRIOR APPLICATION NUMBER: JP 2000-240397

PRIOR FILING DATE: 2000-08-03

PRIOR FILING DATE: 1999-09-27

PRIOR FILING DATE: 1999-09-27

NUMBER OF SEQ ID NOS: 77

SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Masatsugu
APPLICANT: Masatsugu
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APPLICANT: Masatsugu
APPLICANT: Masatsugu
APPLICANT: Masatsugu
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 66501-10543
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-09-27
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR PILING DATE: 2000-08-03
PRIOR PLING DATE: 2000-08-03
PRIOR FILING DATE: 700-08-03
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                    Sequence 69, Application US/10105930 Publication No. US20030009018A1 GENERAL INFORMATION:
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US-10-105-930-72
Sequence 72, Application US/10105930
Publication No. US20030009018A1
GENERAL INFCRMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Homo sapiens
US-10-105-930-69
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US-10-105-930-71
US-10-105-930-69
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APPLICANT: ROSEN et al.

ITILE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REPERENCE: PUZOB
CURRENT APPLICATION UNDBER: US/09/989,442
CURRENT FILING DATE: 2001-11-21
FRIOR PPLICATION NUMBER: 60/179,065
PRIOR PILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: 60/180,628
PRIOR APPLICATION NUMBER: 60/214,886
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR PILING DATE: 2000-08-14
PRIOR PILING DATE: 2000-08-14
PRIOR PILING DATE: 2000-07-25
PRIOR PILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                      Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Miller, Brian S.
APPLICANT: Shetty, Jayarama K.
TITLE OF INVENTION: Modified Forms of Pullulanase
FILE REPERENCE: GC396-2
CURRENT APPLICATION NUMBER: US/09/262,126C
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 9
SOFTWARE: FastSEQ for Windows Version 3.0
        PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR FILING DAFE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 8, Application US/09262126C; Publication No. US20030013180A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 2, Application US/09989442; Publication No. US20030013649A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Unknown US-09-262-126C-8
                                                                                                                                                         ; TYPE: PRT; ORGANISM: Homo sapiens US-10-105-930-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-262-126C-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-989-442-2
                                                                                                                SEQ ID NO 76
LENGTH: 5
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LENGTH: 5
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APPLICANT: Yaguchi, No. US2003009918Aliko
JITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILER REFERENCE: 06501-10548
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
FRIOR APPLICATION NUMBER: DCT/JP00/06654
FRIOR APPLICATION NUMBER: DF 2000-240397
FRIOR FILING DATE: 2000-09-27
FRIOR FILING DATE: 2000-09-27
FRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 75
LENGTH: 5
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APPLICANT: Maeda, Masateugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERBNCE: 06501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 1999-09-27
PRIOR FILING DATE: 1999-09-27
SOFTWARE: PARTSEQ for Windows Version 4.0
SEQ ID NO 74
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; Sequence 76, Application US/10105930
; Publication No. US20030009018A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-105-930-75; Sequence 75, Application US/10105930; Publication No. US20030009018A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                         ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-105-930-74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens
US-10-105-930-75
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Matches 1, Conserv
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R FILING DATE: 2000-08-14

R APPLICATION NUMBER: 60/226,868

R FILING DATE: 2000-08-22

R APPLICATION NUMBER: 60/216,647

R FILING DATE: 2000-07-07

R APPLICATION NUMBER: 60/225,267 APPLICATION NUMBER: 60/236,327 FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/241,785 FILING DATE: 2000-10-20 APPLICATION NUMBER: 60/244,617 APPLICATION NUMBER: 60/236,367 FILING DATE: 2000-09-29 PPLICATION NUMBER: 60/237,039 FILING DATE: 2000-10-02 FILING DATE: 2000-08-14
APPLICATION WUMBER: 60/218,290
ALING DATE: 2000-07-14
APPLICATION NUMBER: 60/225,757 FILING DATE: 2000-07-07
APPLICATION NUMBER: 60/225,270
FILING DATE: 2000-08-14
APPLICATION NUMBER: 60/251,869
FILING DATE: 2000-12-09 APPLICATION NUMBER: 60/236,369 FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/224,519 FILING DATE: 2000-08-14 APPLICATION NUMBER: 60/241,809 FILING DATE: 2000-10-20 APPLICATION NUMBER: 60/249,299 APPLICATION NUMBER: 60/229,343 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/229,345 FILING DATE: 2000-09-01
APPLICATION NUMBER: 60/229,513
ALING DATE: 2000-09-05
APPLICATION NUMBER: 60/231,413 FILING DATE: 2000-09-27 APPLICATION NUMBER: 60/234,274 APPLICATION NUMBER: 60/234,223 FILING DATE: 2000-09-21 FILING DATE: 2000-08-30 APPLICATION NUMBER: 60/224,518 FILING DATE: 2000-08-14 APPLICATION NUMBER: 60/220,964 FILING DATE: 2000-07-26 FILING DATE: 2000-11-01 APPLICATION NUMBER: 60/225,268 60/236,368 FILING DATE: 2000-09-29
APPLICATION NUMBER: 60/251,856 APPLICATION NUMBER: 60/251,868 FILING DATE: 2000-12-08 APPLICATION NUMBER: 60/229,344 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/234,997 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/229,287 FILING DATE: 2000-09-08
APPLICATION NUMBER: 60/229,509 FILING DATE: 2000-10-02 APPLICATION NUMBER: 60/237,038 APPLICATION NUMBER: 60/225,447 APPLICATION NUMBER: 60/216,880 APPLICATION NUMBER: 60/228,924 APPLICATION NUMBER: 60/235,834 FILING DATE: 2000-12-08 FILING DATE: 2000-09-25 FILING DATE: 2000-09-05 FILING DATE: 2000-08-14 FILING DATE: 2000-09-21 FILING DATE: 2000-11-17 FILING DATE: 2000-08-14 APPLICATION NUMBER: PRIOR PRIOR
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R APPLICATION NUMBER: 60/237,037
RR FILING DATE: 2000-10-02
RR APPLICATION NUMBER: 60/237,040
RFILING DATE: 2000-10-02
RR FILING DATE: 2000-10-20
RR FILING DATE: 2000-10-20
RR FILING DATE: 2000-10-20 PILING DATE: 2000-09-14 APPLICATION NUMBER: 60/231,242 FILING DATE: 2000-09-08 PILING DATE: 2000-10-02
APPLICATION NUMBER: 60/236,370
FILING DATE: 2000-09-29
PLING DATE: 2000-10-02
FILING DATE: 2000-10-02 APPLICATION NUMBER: 60/246,474
FILING DATE: 2000-11-08
APPLICATION NUMBER: 60/246,532
APPLICATION NUMBER: 60/246,532
FILING DATE: 2000-11-08 APPLICATION NUMBER: 60/227,182 FILING DATE: 2000-08-22 APPLICATION NUMBER: 60/225,214 FILING DATE: 2000-10-13
APPLICATION WUMBER: 60/239,937
APPLICATION DATE: 2000-10-13
APPLICATION NUMBER: 60/241,787 FILING DATE: 2000-09-06 APPLICATION NUMBER: 60/215,135 FILING DATE: 2000-06-30 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,207 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,245 FILING DATE: 2000-11-17 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,297 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/232,400 APPLICATION NUMBER: 60/249,216 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,210 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/226,681 FILING DATE: 2000-08-22 APPLICATION NUMBER: 60/225,759 APPLICATION NUMBER: 60/225,213 APPLICATION NUMBER: 60/235,836 APPLICATION NUMBER: 60/230,438 APPLICATION NUMBER: 60/225,266 APPLICATION NUMBER: 60/249,218 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,208 APPLICATION NUMBER: 60/249,213 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,212 APPLICATION NUMBER: 60/249,244 APPLICATION NUMBER: 60/249,217 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,211 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,215 APPLICATION NUMBER: 60/249,264 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,214 FILING DATE: 2000-09-27 2000-08-14 FILING DATE: 2000-08-14 FILING DATE: 2000-08-14 FILING DATE: 2000-08-14 FILING DATE: 2000-11-17 2000-11-17 FILING DATE: FILING DATE: FILING DATE: PRIOR PRIOR

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GREATER TRUNGS DEATH STATES AND APPLICANT: Young, Paul APPLICANT: Ruben, Steven M. APPLICANT: Ruben, Steven M. TITLE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation TITLE OF INVENTION: Molecule FILE REFERENCE: PF448
CURRENT APPLICATION NUMBER: US/09/369,248
CURRENT FILING DATE: 1999-08-05
EARLIER APPLICATION NUMBER: 60/073,962
EARLIER APPLICATION NUMBER: 60/073,962
EARLIER APPLICATION NUMBER: 60/078,572
EARLIER FILING DATE: 1998-03-19
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 5
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: Human Neuropeptide Receptor
FILE REFERENCE: PF168P2
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; OTHER INFORMATION: Xaa equals any amino acid
US-09-393-696-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (3)
; OTHER INFORMATION: Xaa equals any amino acid US-09-369-248-5
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CURRENT FILING DATE: 1999-09-10
EARLIER APPLICATION NUMBER: PCT/US95/05616
EARLIER FILING DATE: 1995-05-05
EARLIER FILING DATE: 1995-06-05
MUMBER: US08/462,509
EARLIER FILING DATE: 1995-06-05
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 14
LENGTH: 5
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                                                                                                                               RESULT 71
US-09-369-248-5
Sequence 5, Application US/09369248
Publication No. US20030022276A1
GENERAL INFORMATION:
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US-09-393-696-14
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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ORGANISM: Artificial Sequence
FEATURE:
COMBERING: Alpha5betal integrin-binding peptide
US-10-116-067-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: LINDBOM, Lennart
APPLICANT: LINDBOM, Lennart
APPLICANT: WERR, Joachim
TITLE OF INVENTION: Anti-Inflammatory Medicaments
FILE REFERENCE: 1291-0200P
CURRENT APPLICATION NUMBER: US/10/116,067
CURRENT FILING DATE: 2002-04-05
PRIOR PILING DATE: 1998-10-07
PRIOR FILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: US 60/103,558
PRIOR PILING DATE: 1998-10-07
PRIOR PILING DATE: 1999-10-07
PRIOR FILING DATE: 1999-10-07
NUMBER OF SEQ ID NOS: 5
SOFFWARE: PATENTIN VERSION 3.1
                   PRIOR FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: 60/232,080
PRIOR APPLICATION NUMBER: 60/231,414
PRIOR PILING DATE: 2000-09-08
PRIOR PLING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-14
PRIOR PILING DATE: 2000-09-14
PRIOR PILING DATE: 2000-09-14
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PRIOR PILING DATE: 2000-09-14
PRIOR FILING DATE: 2000-09-14
PRIOR FILING DATE: 2000-09-14
PRIOR PILING DATE: 2000-10-20
PRIOR PILING DATE: 2000-10-8
PRIOR PILING DATE: 2000-10-8
PRIOR PILING DATE: 2000-09-8
PRIOR PILING DATE: 2000-09-8
PRIOR PILING DATE: 2000-09-8
PRIOR PILING DATE: 2000-09-8
APPLICATION NUMBER: 60/232,081
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OTHER INFORMATION: Cytokine receptor extracellular motif found in many species.
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TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR BREAST CANCER
FILE REPERRNCE: A-69959/RMS/DCF
CURRENT APPLICATION NUMBER: US/09/747,377
CURRENT FILING DATE: 2000-12-22
SUFFMER OF SEG ID NOS: 493
SOFTWARE: PatentIn version 3.0
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CURRENT APPLICATION NUMBER: US/09/983,802
CURRENT FILING DATE: 2001-10-25
PRIOR PPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/227,357
PRIOR FILING DATE: EARLIER FILING DATE: 1999-01-08
PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: PCT/US98/13684
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PRIOR FILING DATE: EARLIER FILING DATE: 1996-07-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER PELICATION NUMBER: 60/051,926

PRIOR PILING DATE: EARLIER APPLICATION NUMBER: 60/052,793

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER PILING DATE: 1997-07-08
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (3)..(3) OTHER INFORMATION: "Xaa" at position 3 can be any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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TITLE OF INVENTION: 123 Human Secreted Proteins
                                                                                                                                                                                         Sequence 493, Application US/09747377
Publication No. US20030022255A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09983802
Publication No. US20030022185A1
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GENERAL INFORMATION:
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Best Local Similarity
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                                                                                                                                                    US-09-747-377-493
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OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino OTHER INFORMATION: acids
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; Publication No. US20030022327A1
; Publication No. US20030022327A1
; GENEAL INFORMATION:
; APPLICANT: Ruben et al.
; TITLE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation
; TITLE OF INVENTION: Molecule
; FILE REFERENCE: PF448P2
; CURRENT APPLICATION NUMBER: US/10/062,523
PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR APPLICATION NUMBER: EARLIER PAPPLICATION NUMBER: 60/051,919

PRIOR APPLICATION NUMBER: EARLIER PAPPLICATION NUMBER: 60/051,928

PRIOR APPLICATION NUMBER: EARLIER PAPPLICATION NUMBER: 60/055,722

PRIOR APPLICATION NUMBER: EARLIER PAPPLICATION NUMBER: 60/055,723

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,723

PRIOR PILING DATE: EARLIER FILING DATE: 1997-09-18

PRIOR PILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR PELING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR PILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,949

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PRIOR APPLICATION NUMBER: EARLIE
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SOFTWARE: Patentin Ver. 2.0
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Perfect score:

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Sequence 2, Appli
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Sequence 13, Appl
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Sequence 11, Appl
Sequence 11, Appl
Sequence 100, App
Sequence 454, App
Sequence 454, App
Sequence 228, App
Sequence 14, Appl
Sequence 15, Appl
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6 US-09-214-711-2

6 US-09-214-711-3

10 US-09-603-440A-1

10 US-09-507-361-3

10 US-09-957-0986-725

10 US-09-957-0986-725

10 US-09-957-0986-725

10 US-09-957-0986-725

10 US-08-21257-11

10 US-09-21257-11

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10 US-09-2125-14

10 US-09-2125-14

10 US-09-515-965A-19-1

10 US-09-515-965A-19-1

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10 US-09-612-210-21

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US-10-197-954-139
US-10-200-923-8
US-10-200-923-10
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Sequence 114, App
Sequence 11, Appl
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Sequence 114, Appl
Sequence 111, Appl
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19: /cgm2_6/ptodata/1/paa/USO99_COMB.pep:*
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19: /cgm2_6/ptodata/1/paa/USO99_COMB.pep:*
10: /cgm
                     GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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PCT-US99-26277-114
US-09-601-558-11
US-09-811-047B-114
US-09-891-047C-114
US-09-894-594-11
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Maximum Match 100%
Listing first 100 summaries
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43, Appl 51, Appl 30, Appl 1, Appl 582, App 791, App 791, App

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Sequence 99, Appl
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Sequence 48, Appl
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                                                     Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Magdalena
APPLICANT: Blaszczyk-Thurin, Magdalena
APPLICANT: Reiber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer
CURRENT APPLICATION NUMBER: PCT/US99/26277
CURRENT FILING DATE: 1999-11-05
EARLIER APPLICATION NUMBER: 60/107,478
EARLIER FILING DATE: 1999-11-06
NUMBER OF ERQ ID NOS: 121
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Trustees of the University of Pennsylvania TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens FILE REFERENCE: upn3544 CURRENT APPLICATION NUMBER: PCT/US99/02405 CURRENT FILING DATE: 1999-02-04 EARLIER APPLICATION NUMBER: 60/073,690 EARLIER PILING DATE: 1998-02-04 NUMBER: PELING DATE: 1998-02-04 SOFTWARE: PALENT NOS: 60 SOFTWARE: PALENT NOS: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 87.0%; Score 20; DB 1; Length 5; Best Local Similarity 60.0%; Pred. No. 4.2e+06; Matches 3; Conservative 0; Mismatches 2; Indels
S US-10-200-923-11
S US-10-200-923-12
S US-10-200-923-13
S US-10-237-673-13
7 US-60-366-75-65
US-60-366-75-65
US-09-392-941-83
7 US-09-392-941A-169
9 US-09-392-941A-169
9 US-09-392-941A-169
PCT-US00-07724-75
PCT-US00-07724-75
PCT-US02-14843-4
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PCT-US96-13921-11
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APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blasczyk-Thurin, Magdalena
APPLICANT: Blasczyk-Thurin, Magdalena
APPLICANT: Blasczyk-Thurin, Magdalena
TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer
FILE REPERENCE: MSY93ADSA
CURRENT APPLICATION NUMBER: US/09/831,047B
CURRENT FILING DATE: 2001-08-14
FRICA APPLICATION NUMBER: 60/107,478
FRICA APPLICATION NUMBER: 60/107,478
FRICA FILING DATE: 1998-11-06
NUMBER OF SEQ ID NOS: 121
SCOTWARE PATENTIN Ver. 2.0
SEQ ID NO 114
LENGTH: 5
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; GENERAL INFORMATION:
; APPLICANT: Kieber-Emmons, Thomas
; APPLICANT: The Trustees of the University of Pennsylvania
TITLE OF INVENTION: Peptide Mimotopes Of Carbohydrate Antigens; FILE REFERENCE: UR3857
; CURRENT APPLICATION NUMBER: US/09/601,558
; CURRENT FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: 60/073,690
; PRIOR PILING DATE: 1999-02-04
; PRIOR PILING DATE: 1998-02-04
; RIOR PILING DATE: 1998-02-04
; NUMBER OF SEQ ID NOS: 60
; SOFFWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                            87.0%; Score 20; DB 1; Length 5; 60.0%; Pred. No. 4.2e+06; Live 0; Mismatches 2; Indels
                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: peptido-mimetic of a Lewis antigen
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                         TYPE: PRT ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 60.03
Matches 3; Conservative
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Best Local Similarity 60.09
Matches 3; Conservative
                                                                                                          ) OTHER INFORMATION OTHER INFORMATION PCT-US99-26277-114
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US-09-601-558-11
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LENGTH: 5
LENGIH: 5
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TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR TITLE OF INVENTION: ACTIVITY ACTIVITY ACTIVITY ACTIVITY OF INVENTION: ACTIVITY ACT
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APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TILE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
                          87.0%; Score 20; DB 22; Length 5; 60.0%; Pred. No. 4.2e+06; ive 0; Mismatches 2; Indels
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OTHER INFORMATION: Xaa is any amino acid
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GENERAL INPORMATION:
APPLICANT: GUY, RODNEY
APPLICANT: DARIMONT, BEATRICE
APPLICANT: DARIMONT, BEATRICE
APPLICANT: PENG, WEIJUN BAPLICANT:
APPLICANT: PENG, WEIJUN BAPLICANT: PENG, WEIJUN BAPLICANT: PENG, WEIJUN BAPLICANT: APPLICANT: PENG, WEIJUN BAPLICANT: APPLICANT: APAMONO, KETTH
APPLICANT: YAWAMOTO, KETTH
APPLICANT: ARAMONO, KETTH
APPLICANT: ARAMONO, JAMES
APPLICANT: KUNTZ, IRWIN
APPLICANT: KUNTZ, IRWIN
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Bottger, Volker
Bottger, Angelica
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Matches 5; Conservative
                                                                                                               3; Conservative
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                                                                       Best Local Similarity
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APPLICANT: Lane, Da
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US-09-214-371-2
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APPLICANT:
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APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Wonzard: Karbassi, Behjatolah
ITILE OP INVENTION: Same
FILE OP INVENTION: Same
FILE OF INVENTION: Same
FILE OP INVENTION NUMBER: US/09/894,594
FILE OF TILING DATE: 2000-11-07
FILE OP INVENTION NUMBER: E0/073,690
FILE OF TILING DATE: 1998-02-04
FILE OF TILING DATE: 1998-02-04
FILE OF TILING DATE: 2000-06-28
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; GENERAL INFORMATION:
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Magdalena
APPLICANT: Kieber-Emmons, Thomas
TILLE REPERBNCE: WST93ABGSA
CURRENT APPLICATION NUMBER: US/09/831,047C
CURRENT APPLICATION NUMBER: 08/09/831,047C
CURRENT FILING DATE: 2001-05-03
PRIOR PLICATION NUMBER: 60/107,478
PRIOR PLICATION NUMBER: 60/107,478
PRIOR PLILIG DATE: 1998-11-06
NUMBER OF SEQ ID NOS: 121
SOUTHWER PAPELICATION 3.1
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                                                                                         Score 20; DB 22; Length 5;
Pred. No. 4.2e+06;
0: Mismatches 2; Indels
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                                                                                                  Query Match
87.0%; Score 20; DB
Best Local Similarity 60.0%; Pred. No. 4.2e
Matches 3; Conservative 0; Mismatches
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OTHER INFORMATION: Novel Sequence
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Best Local Similarity
Matches 3; Conserv
US-09-831-047B-114
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US-09-831-047C-114
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LENGTH: 5
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CURRENT FILING DATE: 2000-01-19
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SOFTWARE: Patentin version 3.0
SEQ ID NO 3
LENGTH: 5
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                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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Matches 5; Conservative
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LENGTH: 5
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JOURNELL IN WARTHION:

APPLICANT: Baxter, John D.

APPLICANT: Baxter, John D.

APPLICANT: Feder J.

APPLICANT: Fletterick, Robert J.

APPLICANT: Weather, Peter J.

APPLICANT: Wagner, Richard L.

APPLICANT: Wagner, Richard L.

APPLICANT: Wammoto, Keith R.

TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR

TITLE OF INVENTION: COACTIVATOR BINDING

FILE REPRENCE: UCAL-253/02US

CURRENT FILING DATE: 1999-03-30

CURRENT FILING DATE: 1999-03-30

EARLIER APPLICATION NUMBER: US 60/079,956

EARLIER PILING DATE: 1998-03-30

NUMBER OF SEQ ID NOS: 51

SOFTWARE PATEUTING DATE: 2.1
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TITLE OF INVENTION: MATERIALS AND METHODS RELATING TO
TITLE OF INVENTION: INHIBITING THE INTERACTION OF p53 AND MDM2
FILE REFERENCE: MEMBERS: 001APC
CURRENT APPLACATION NUMBER: US/09/403,440A
                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence:peptide OTHER INFORMATION: Where Xaa may be any amino acid
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                                                                                                                                                                                                                                                                                                                                   Length 5;
                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                Query Match 82.6%; Score 19; DB 16; L. Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 5; Conservative 0; Mismatches 0;
CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 PRIOR APPLICATION NUMBER: PCT/EP97/03549 PRIOR FILING DATE: 1997-07-04 NUMBER OF SEQ ID NOS: 83 SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: SITE
LOCATION: (2)..(3)
OTHER INFORMATION: Xaa = Any Amino Acid
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1 FXXLW 5
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                                                                                                              SOFTWARE: 1
SEQ ID NO 2
LENGTH: 5
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Matches
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APPLICANT: RICHARD, WAGER
APPLICANT: BILAN, WEST
APPLICANT: BILAN, WEST
APPLICANT: BILAN, WEST
APPLICANT: AND BILAN, WEST
APPLICANT: ARNOLD, JAMES
APPLICANT: WINTZ, IRWIN
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR ACTIVITY
FILE REFERENCE: 981-016-999
CURRENT PILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: US 60/079,965
PRIOR PRIOR APPLICATION NUMBER: US 60/113,146
PRIOR FILING DATE: 1999-03-30
PRIOR FILING DATE: 1999-03-30
PRIOR FILING DATE: 1999-03-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                FEATURE: OTHER INFORMATION: This represents a consensus sequence representing OTHER INFORMATION: humans and mice.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 82.6%; Score 19; DB 18; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 5; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: PCT/GB98/01144
PRIOR FILING DATE: 1998-04-20
PRIOR APPLICATION NUMBER: GB 9708092.3
PRIOR FILING DATE: 1997-04-22
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Xaa = any amino acid US-09-403-440A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/09609361; GENERAL INFORMATION:
APPLICANT: GUY, RODNEY
APPLICANT: BAXTER, JOHN
APPLICANT: DAXTER, JOHN
APPLICANT: FENG, WEIJUN
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NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 5
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ORGANISM: Artificial Sequence
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Matches 2; Conservative
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ORGANISM: Homo Sapien
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PCT-US02-23257-10
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LENGTH: 5
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Sequence 226, Application US/09957806A
; Sequence 226, Application US/09957806A
; GENERAL INPORMATION:
; APPLICANT: ROGGEN, Erwin
; APPLICANT: Syendeen, Allan
; APPLICANT: Syendeen, Allan
; APPLICANT: Friis, Eaben
; APPLICANT: Osten, Claus
; TITLE OF INVENTION: PROTEIN VARIANTS HAVING MODIFIED IMMUNOCENICITY
; FILE REFRENCE: 10021.204-US
; CURRENT APPLICATION UNBER: US/09/957,806A
; CURRENT FILING DATE: 2001-09-21
; NUMBER OF SEQ ID NOS: 248
; SOFTWARE: Patentin version 3.1
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Sequence 791, Application PC/TUS0103988
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
CURRENT APPLICATION NUMBER: PCT/US01/03988
CURRENT PILING DATE: 2001-02-07
PRIOR APPLICATION NUMBER: 09/515,965
PRIOR APPLICATION NUMBER: 09/515,965
PRIOR APPLICATION NUMBER: 09/515,965
PRIOR SEQ ID NOS: 1589

NUMBER OF SEQ ID NOS: 1589
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82.6%; Score 19; DB 23; Length 5;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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APPLICANT: Koister, Hubert
APPLICANT: Siddigi, Suhaib
APPLICANT: Little, Daniel
TITLE OF INVENTION: Capture Compounds, Collections Thereof
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65.2%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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; GENERAL INFORMATION:
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TITLE OF INVENTION: And Methods For Analyzing The Proteome And Complex TITLE OF INVENTION: Compositions FILE REPRENCE: 24743-2305
CURRENT APPLICATION NUMBER: PCT/US02/22821
CURRENT PILING DATE: 2002-07-16
PRIOR APPLICATION NUMBER: 60/306,019
PRIOR FILING DATE: 2001-07-16
PRIOR APPLICATION NUMBER: 60/306,113
PRIOR FILING DATE: 2001-08-21
PRIOR APPLICATION NUMBER: 60/363,433
PRIOR FILING DATE: 2002-03-11
NUMBER OF SEQ ID NOS: 149
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
APPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REPERENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/307,228
MINDER OF THING DATE: 2001-07-23
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GENERAL INFORMATION:
APPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REFERENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/307,228
PRIOR FILING DATE: 2001-07-23
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SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2, Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REFERENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT APPLICATION NUMBER: US 60/307,228
PRIOR FILING DATE: 2001-07-23
PRIOR FILING DATE: 2001-07-23
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                              Query Match
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11, Application PC/TUS0223257
GENERAL INFORMATION:
FAPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: ANALGESIC METHODS
FILE REPREBENCE: S0033/0.10002
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT APPLICATION NUMBER: US 60/307,228
FRIOR APPLICATION NUMBER: US 60/307,228
FRIOR FILING DATE: 2001-07-23
NUMBER OF SEQ ID NOS: 14
SOSTWARE: FRSESQ FOR Windows Version 4.0
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Sequence 12, Application PC/TUS0223257
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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PCT-US02-23257-12
                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Synthetic PCT-US02-23257-11
                                                                                            ; FEATURE:
; OTHER INFORMATION: Synthetic
PCT-US02-23257-10
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PCT-US02-23257-11
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; SEQ ID NO 10
                        LENGTH: 5
                                            TYPE: PRT
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Sequence 1, Application PC/TUS9815227
(GENERAL INFORMATION:
APPLICANT: The New York Blood Center, Inc.
TITLE OF INVENTION: MONOSPECIFIC ANTIBODY REACTIVE WITH
TITLE OF INVENTION: MATRIX METALLOPROTEINASE CLEAVAGE PRODUCTS OF FIBRIN (OGEN)
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                           JAPERANT: THE BLY, AND WOMEN'S HOSPITAL, INC.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REPERENCE: 50073/010002
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT APPLICATION NUMBER: DCT/US02/23257
PRIOR APPLICATION NUMBER: US 60/307,228
PRIOR FILING DATE: 201-07-23
NUMBER OF FILING DATE: 201-07-23
SOFTWARE: FRASEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diskette, 3.50 inch, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ), OTHER INFORMATION: Ile at position 3 is alle PCT-USO2-23257-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               454-16 PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US98/15227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Hoffmann & Baron, LLP STREET: 350 Jericho Turnpike
                                                                                                                                                  ; Sequence 13, Application PC/TUS0223257; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/900,895
FILING DATE: 7/25/97
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Baron, Ronald J.
REGISTRATION NUMBER: 29,281
REFERENCE/DOCKET NUMBER: 454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IBM compatible
SYSTEM: MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WordPerfect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jericho
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                                                                                                                                 PCT-US02-23257-13
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LENGTH: 5
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4 LW 5
                                        4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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GENERAL INFORMATION:
APPLICANT: TRIMERIS, INC.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-052-228
CURRENT APPLICATION NUMBER: PCT/US99/11219
CURRENT PILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 1542
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                   ;
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ZIP: 20005

COMPUTER READABLE FORM:

COMPUTER: Floppy disk

COMPUTER: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/585,141

TING DATE: 19901009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11, Application US/07585141
GENERAL INFORMATION:
APPLICANT: Schramm, Wolfgang
APPLICANT: Schramm, Hans J.
TITLE OF INVENTION: Agent For Inhibiting Symmetrical
TITLE OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
65.2%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PEATURE:
COTHER INFORMATION: Core polypeptide
PCT-US99-11219-791
  TELEPHONE: (516) 822-3550
TELEFAX: (516) 822-3582
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                       ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US98-15227-1
                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 1201 New CITY: Washington STATE: DC
TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                           PCT-US99-11219-791
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LENGTH: 5
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Sequence 17, Application US/07608569;
GENERAL INFORMATION:
APPLICANT: Tainer, John A
APPLICANT: Ruhn, Leslie
APPLICANT: Boissinot, Maurice
APPLICANT: Pisher, Cindy
APPLICANT: Parge, Hans B
APPLICANT: Griffin, John H
APPLICANT: Mullenbach, Guy T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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ADDRESSEE: The Scripps Research Institute, Office of ADDRESSEE: Patent Counsel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: Eloppy disk
COMPUTER: Elem PC Compatible
OPERATING SYSTEM: PC-005/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/07/608,569
FILING DATE: 19901102
CLASSIFICATION 1424
PRIOR APPLICATION NUMBER: US 07/608539
FILING DATE: 01-NOV-1990
ATTORNEY/AGENT INFORMATION:
ANALYSEE AFFER THE TRANSANTION:
ANALYSEE AFFER THE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 65.2%; Score 15; DB 3; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/EP90/00219
FILING DATE: 09-FEB-1990
PRIOR APPLICATION DATA:

APPLICATION NUMBER: DE P 39 04 040.2
FILING DATE: 10-FEB-1989
ATTORNEY/AGENT INFORMATION:

NAME: Ihnen, Jeffrey L.

REGISTRATION NUMBER: 28957
REFERENCE/POCKET NUMBER: 18644-96040
TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEPAK: 202-962-8300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: CA
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REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: SCR0348P
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-544-2937
TELEFAX: 619-554-6312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 5 amino acids TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide; HYPOTHETICAL: YES
US-07-585-141-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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US-08-344-185B-454
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: MUERHOFF, ANTHONY T.
APPLICANT: MUERHOFF, ANTHONY T.
APPLICANT: BUIJK, S.L.
APPLICANT: BUIJK, S.L.
APPLICANT: MUSHAHWAR, ISA K,
TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-D, NON-E
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 539
CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                                                                              65.2%; Score 15; DB 3; Length 5; ilarity 100.0%; Pred. No. 4.2e+06; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,185B
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 100, Application US/08344185B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PILOT-MATTAS, T.J.
DAWSON, GEORGE J.
SCHLAUDER, GEORGE G.
DESAI, SURESH M.
LEARY, THOMAS P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: SIMONS, JOHN N.
      TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                       SEQUENCE CHARACTERISTICS:
  INFORMATION FOR SEQ ID NO:
                                                                                                MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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CLASSIFICATION: 435
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Matches 2; Conserv
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RESULT 25

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GENERAL INFORMATION:

APPLICANT: SIMONS, JOHN N.

APPLICANT: DAWSON, GEORGE J.

APPLICANT: BARSON, GEORGE G.

APPLICANT: SCHLAUDER, GEORGE G.

APPLICANT: DESAI, SUREAH

APPLICANT: LEARY, THOMAS P.

APPLICANT: BUIJK, S.L.

APPLICANT: MUSHAPHWAR, ISA K.

TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-D, NON-E

TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE

NUMBER OF SEQUENCES: 539

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                          APPLICANT: DESAI, SURESH M.
APPLICANT: LEARY THOMAS P.
APPLICANT: MUBHOFF, ANTHONY T.
APPLICANT: BUIJK, S.L.
APPLICANT: MUSHAWAR, ISA K.
TITLE OF INVENTION: MON-A, NON-B, NON-C, NON-D, NON-E
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 65.2%; Score 15; DB 7; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 539
CORRESPONDENCE ADDRESS:
ADDRESSE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/344,185B
; Sequence 454, Application US/08344185B ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/08344185C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: POREMISKI, PRISCILLA E.
TELECOMMINICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                      APPLICANT: SIMONS, JOHN N.
APPLICANT: PLIOT-MATILAS, T.J.
APPLICANT: DAWSON, GEORGE J.
APPLICANT: SCHLAUDER, GEORGE G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-344-185B-454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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US-08-344-185C-100
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APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
NUMBER OF SEQUENCES: 116
CORRESPONDENCE ADDRESS:
                                                                                                                                                     Query Match 65.2%; Score 15; DB 7; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA

ZIP: 60064-3500

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550A
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 65.2%; Score 15; DB 8; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SSEE: ABBOTT LABORATORIES D377/AP6D
1: 100 ABBOTT PARK ROAD
ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                             RESULT 28
US-08-424-550A-228
IS-08-424-550A-228
Sequence 228, Application US/08424550A
GENERAL INFORMATION:
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: THOMAS P. LEARY
APPLICANT: THOMAS P. LEARY
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
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ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 55:
TELECOMMUNICATION INFORMATION:
TELECHONE: 708-937-6365
TELEFAX: 708-938-2623
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INFORMATION FOR SEQ ID NO: 228:
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                 LENGTH: 5 amino acids
                                  ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-344-185C-454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-424-550A-228
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US-08-424-550B-228
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APPLICANT: BILOT-MATIAS, T.J.
APPLICANT: BLOT-MATIAS, T.J.
APPLICANT: SCHLAUDER, GEORGE J.
APPLICANT: SCHLAUDER, GEORGE G.
APPLICANT: LEARY, THOMAS P.
APPLICANT: LEARY, THOMAS P.
APPLICANT: BUICK, ANTHONY T.
APPLICANT: WUSENGYF, ANTHONY T.
APPLICANT: WUSENGYF, ANTHONY T.
APPLICANT: WUSENGY S.L.
APPLICANT: WUSENGY S.L.
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE CORRESPONDENCE ADDRESSES:
ADDRESSEE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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  STÄTE: IL
COUNTRY: USA
ZIP: 60064-3500
COMPUTER REABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,185C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,185C
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 454, Application US/08344185C GENERAL INFORMATION:
                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
TELEFAX: 708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: POREMBEKI, PRISCILLA E.
TELECOMMINICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     454:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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IL
USA
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                                                                                                                                                                                                                                                    APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
ADDRESSEE: ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 47, Application US/08432147
GENERAL INFORMATION:
APPLICANT: Kishore, Ganesh M.
APPLICANT: Arbondel, Vincent J.
APPLICANT: Arbondel, Vincent J.
APPLICANT: Gomerville, Christopher R.
APPLICANT: Gomerville, Christopher R.
TITLE OF INVENTION: Altered Linolenic and Linoleic Acid
TITLE OF INVENTION: Content in Plants
NUMBER OF SEQUENCES: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           B: Lawrence M. Lavin, Jr., Monsanto Co., BB4F 700 Chesterfield Parkway North
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:

ATCNREY/AGBNT INFORMATION:
NAME: PORENBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEFAX: 708-938-2623
                                SENERAL INFVENCEALLOW:
APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURESH M. DESAI
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUBRHOFF
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: STARM
Sequence 228, Application US/08424550B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                : 5 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS: ADDRESSEE: Lawrence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 700 Chest
CITY: St. Louis
STATE: Missouri
COUNTRY: USA
                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-424-550B-228
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US-08-432-147-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 65.2%; Score 15; DB 8; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/467,344A
FILING DATE: 07-Jun-1995
CLASSIFICATION: <UNKnown>
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ADDRESSEE: ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
OPENATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/432,147
FILING DATE: 05-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA

APPLICATION NUMBER: PCTUS94/01321

FILING DATE: 04-FEB-1994

PRIOR APPLICATION NUMBER: EP 94908724.1

FILING DATE: 04-FEB-1994

ATTORNEY/AGENT INFORMATION:
NAME: Lavin Jr., Lawrence M.
REGISTRATION NUMBER: 30,768

REFERENCE/COCKET NUMBER: 30,768

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
GENERAL INFORMATION:
HAPLICANT: JOHN N. SIMONS
GEORGE J. DAWSON
GEORGE J. DAWSON
GEORGE J. DAWSON
HAPPING STREET
THOMAS P. LEARY
JAMES C. ERKER
SHERI L. BUIJK
HAPPING STREET
HOMAS P. LEARY
HOMAS 
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (314)537-6047
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 716
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: peptide US-08-432-147-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                               FILING DATE: 05 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 LW 5
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COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: PORENBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 708-22.
TELEPHONE: 708-238-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
                         ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-486-749-228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , MOLECULE TYPE: protein US-08-488-445A-228
      amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 34
US-08-488-446-228
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APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWGON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURESH M. DESAI
APPLICANT: THOMAS P. LERAY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. ERKER
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUJJK
APPLICANT: SHERI L. BUJJK
APPLICANT: STERI L. BUJJK
APPLICANT: TSA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-D, NON-E HEPATITIS
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 65.2%; Score 15; DB 8; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PARTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,749
                                  ATTORNEY/AGENT INFORMATION:

ATTORNEY/AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-938-5623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: ABBOTT LABORATORIES D377/AP6D STREET: 100 ABBOTT PARK ROAD CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 228:
US-08-467-344A-228
*PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/424,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               33,207
ER: 5527.PC.01
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 228, Application US/08486749 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/POCKET NUMBER: 5527.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-931-6365
TELEPRAK: 708-938-2623
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                      Gaps
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TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEFATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
CORRESPONDENCES: 716
CORRESPONDENCE ADDRESS:
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Query Match 65.2%; Score 15; DB 8; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,445A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 65.2%; Score 15; DB 8; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGULT 33
US-08-488-445A-28
IS-08-488-445A-28
Sequence 228, Application US/08488445A
Sequence 228, Application US/08488445A
Sequence 228, Application US/08488445A
Sequence 228, Application US/08488445A
APPLICANT: JOHN N. SIMONS
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. BEKER
APPLICANT: SHERI L. BUIJK
APPLI
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
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FILING DATE:
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                                                                                    COMPUTER:
    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 LW 5
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GENERAL INFORMATION:
APPLICANT: HIGGER
APPLICANT: BINKELBORG, LUDGER
APPLICANT: KRAMP, WOLGANG
APPLICANT: SCHIER, HANS-MARTIN
TITLE OF INVENTION: CHELATING AGENTS OF THE TYPE XNISIOI FOR RADIOACTIVE I
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLIESLER, DUBB, MEYER & LOVEJOY
STRRET: FOUR Embarcadero Center, Suite 400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                 APPLICANT: JOHN N. SIMONS
APPLICANT: JOHN N. SIMONS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: STRESH M. DESAI
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT WUERHOFF
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUIJOK
APPLICANT: ISA K. MUSHAHARA
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC_compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,446
                                                                                                                                                                                                                                                                                                                                 SEE: ABBOTT LABORATORIES D377/AP6D: 100 ABBOTT PARK ROAD ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: POREMEST, PRISCILLA E.

REGISTRATION NUMBER: 33,207

REFERENCE/DOCKET NUMBER: 5527.PC.01

TELECOMMUNICATION INFORMATION:

TELEPAX: 708-938-2635

INFORMATION FOR SEQ ID NO: 228:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
Application US/08488446
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-488-446-228
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Best Local Similarity 1
                                                                                                                                                                                                                                                                                                                                                                                                                                          60064-3500
; Sequence 228, Applica ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
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US-08-530-325-14
                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                          STREET:
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RESULT 36
US-08-530-127-14

Sequence 14, Application US/08530327
Sequence 14. Application US/08530327
Sequence 14. Application US/08530327
Sequence 14. Application US/08530327
Sequence 14. Applicant: ERBER, SEBASTIAN
APPLICANT: BINKELEDROR, LUDGER
APPLICANT: NOLL, BERNHARD
APPLICANT: NOLL, BERNHARD
APPLICANT: NOLL, BERNHARD
APPLICANT: NOLL, BERNHARD
CORRESPONDENCES: 16
NUMBER OF SEQUENCES: 16
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSES: FLIESLER, DUBB, MEYER & LOVEJOY
STREET: Four Embarcadero Center, Suite 400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: WBLT 2005 (WO) PA TELECOMMUNICATION INFORMATION: TELEPHONE: 415-362-3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUFTWARE: LESTERS DOS
SUFTWARE: FastSEO Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/530,327
FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                            CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00371
FILING DATE: 29-MAR-1994
APPLICATION NUMBER: P 43 11 023.1
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAMME: Dubb, Hubert E
REGISTRATION NUMBER: 25,329
                                                                                        SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/530,325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                             IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-362-2928
                            Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide HYPOTHETICAL: NO
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FRAGMENT TYPE: i
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OTHER INFORMATION: Sequence of a peptide synthesized in Examples 11 and 12 from L-fo OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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                           APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Azitomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REPERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122A
CURRENT FILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: D7 -176904
EARLIER PILING DATE: 1995-06-21
EARLIER PILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 38
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 65.2%; Score 15; DB 13; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: UP 7-176904
EARLIER FILING DATE: 1996-06-21
EARLIER FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SEQ ID NO 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 43, Application US/08981122A
; GENERAL INFORMATION:
; APPLICANT: Hatanaka, Yoshihiro
                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: AMIDATION LOCATION: 5
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         GENERAL INFORMATION:
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OTHER INFORMATION: Sequence of a peptide synthesized in Example 2 from L-form
OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
OTHER INFORMATION: synthesizing system (RaMPS)
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APPLICANT: Aritomi, Masaharu
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILIS REFERENCE:
CURRENT PILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: UP 7-176904
EARLIER PILING DATE: 1995-06-21
NUMBER OF FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65.2%; Score 15; DB 9; Length 5; 100.0%; Pred. No. 4.2e+06; vative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                WBLT 2007 (WO) PA
                                          FILING DATE: 29-MRR-1994 APPLING DATE: 29-MR-1994 FILING DATE: 21-MR-1993 ATTORNEY/AGENT INFORMATION: NAME: Dubb, Hubert E REGISTRATION NUMBER: 25,329 REFERENCE/DOCKET NUMBER: WBLT 2007 (
                           PCT/DE94/00369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 37
US-08-981-122-15
Sequence 15, Application US/08981122A
; GENERAL INFORMATION:
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US-08-981-122-38
; Sequence 38, Application US/08981122A
                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-362-3800
                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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Best Local Similarity
Matches 2; Conserv
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US-08-981-122-15
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APPLICANT: Barney, S.
APPLICANT: Guthrie, K.
APPLICANT: Merutka, G.
APPLICANT: Anwer, M.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC PROPERTIES
GENERAL INFORMATION:
APPLICANT: Genencor International, Inc.
TITLE OF INVENTION: Novel EGIII-Like Enzymes, DNA Encoding
TITLE OF INVENTION: Novel EGIII-Like Enzymes, DNA Encoding
TITLE OF INVENTION: Such Enzymes and Methods for Producing Such Enzymes
FILE REFERENCE: GCS16-2-PCT
CURRENT APPLICATION NUMBER: US/09/284,327
CURRENT APPLICATION DATE: 1999-04-10
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FASTERQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Novel Prokaryotic Polynucleotides,
TITLE OF INVENTION: Polypeptides and Their Uses
FILE REFERENCE: P50596
CURRENT APPLICATION NUMBER: US/09/308,823A
CURRENT FILING DATE: 1999-05-25
NUMBER OF SEQ ID NOS: 661
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 65.2%; Score 15; DB 16; L
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/09/350,641

CURRENT FILING DATE: 1999-07-09

PRIOR APPLICATION NUMBER: 09/315,304

PRIOR FILING DATE: 1999-05-20

PRIOR APPLICATION NUMBER: 09/082,279

PRIOR FILING DATE: 1998-05-20

NUMBER OF SEQ ID NOS: 1589

SOFTWARE: FASTSEQ FOr Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: PRT; ORGANISM: Staphylococcus aureus
US-09-308-823A-582
                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 43
US-09-308-823A-582
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US-09-350-641-791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid CTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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TITLE OF INVENTION: The Use of Consensus Sequences for Targeted Homologous
TITLE OF INVENTION: Gene Isolation and Recombination in Gene Families
FILE REFERENCE: A-65678-1/RFT/NBC
CURRENT APPLICATION NUMBER: U5/09/209,676
CURRENT FILING DATE: 2002-06-18
PRIOR APPLICATION NUMBER: U5 60/070,734
PRIOR FILING DATE: 1997-12-11
NUMBER OF SEQ ID NOS: 95
SOFTWARE: Patentin Ver. 2.0
                                                                                                        APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
IITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
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US-09-209-676-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 13; Length 5; ilarity 100.0%; Pred. No. 4.2e+06; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                  FILE REPERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122A
CURRENT FILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: UP 7-176904
EARLIER FILING DATE: 1995-06-21
EARLIER FILING DATE: 1995-06-21
EARLIER FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
                                                    Sequence 51, Application US/08981122A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 30, Application US/09209676; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Ver. 2.0
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Matches 2; Conserva
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US-09-209-676-30
                                   US-08-981-122-51
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FEATURE:
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SEQ ID NO 791
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LENGTH: 5
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APPLICANT: Anwer, M.
APPLICANT: Lambert, D.
TITLE OP INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OP INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
CURRENT APPLICATION NUMBER: US/09/350,641B
PRIOR PILING DATE: 1999-01-09
PRIOR PELICATION NUMBER: 09/315,304
PRIOR PLING DATE: 1999-05-20
PRIOR PLING DATE: 1998-05-20
PRIOR PLING DATE: 1998-05-20
PRIOR PELICATION NUMBER: 09/082,279
NUMBER OF SEQ ID NOS: 1757
SOFTWARE: FastSEQ for Windows Version 3.0
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GENERAL INFORMATION:
APPLICANT: Barrey, S.
APPLICANT: BARREY, S.
APPLICANT: GLICHIE, K.
APPLICANT: ADMENT OF THE SET OF THE SE
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                                                                                                                                                ; OTHER INFORMATION: Core polypeptide
US-09-350-641-791
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US-09-350-6418-791
LENGTH: 5
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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ORGANISM: Artificial Sequence
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GENERAL INFORMATION:
APPLICANT: Probst, Peter
APPLICANT: Bhatia, Ajay
APPLICANT: Skeiky, Yasir
APPLICANT: Skeiky, Yasir
APPLICANT: Fang, Hang
APPLICANT: Stromberg, Erica Jean
APPLICANT: Stromberg, Erica Jean
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121.469C2
CURRENT FILING DATE: 1999-10-01
NUMBER OF SEQ ID NOS: 109
NUMBER OF SEQ ID NOS: 109
SOFTMARE: FASESEQ for Windows Version 3.0
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APPLICANT: Bhatia, Ajay
APPLICANT: Bhatia, Ajay
APPLICANT: Eling, Steve
APPLICANT: Filing, Steve
APPLICANT: Fang, Hang
APPLICANT: Gen, Shyian
APPLICANT: Stromberg, Erica Jean
APPLICANT: Enghart, Susan E.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INPECTION
                                                                                                                                             65.2%; Score 15; DB 17; Length 5; 100.0%; Pred. No. 4.2e+06; tive 0; Mismatches 0; Indels
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65.2%; Score 15; DB 18; L
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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CURRENT FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 168
SOFTWARE: FastSEQ for Windows Version 3.0
                                              FEATURE:

COTHER INFORMATION: Core polypeptide
US-09-350-641C-791
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US-09-426-571-9
Sequence 9, Application US/09426571
GENERAL INFORMATION:
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ORGANISM: Chlamydia trachomatis
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 2; Conservative
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Query Match
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APPLICANT: Ping, Stewe, Jeff
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121-469C4
CURRENT FILING DATE: 1999-12-03
NUMBER OF SEQ ID NOS: 310
SOFTWARE: PastSEQ for Windows Version 3.0/4.0
SEQ ID NO 9
LENGTH: 5
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GENERAL INFORMATION:
APPLICANT: Probst, Peter
APPLICANT: Skeiky, Yasir
APPLICANT: Skeiky, Yasir
APPLICANT: Brain, Steve
APPLICANT: Maisonneuve, Jeff
APPLICANT: Maisonneuve, Jeff
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 65.2%; Score 15; DB 18; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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                                    h Similarity 100.0%; Pred. No. 4.2e+06; 2; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION WUMBER: US/09/454,684
CURRENT FILING DATE: 1999-12-03
NUMBER OF SEQ ID NOS: 303
SOFTWARE: FastSEQ for Windows Version 3.0/4.0
SEQ ID NO 9
LENOTH: 5
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Best Local Similarity
Matches 2; Conserv
                                         Query Match
Best Local Similarity
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US-09-454-684A-9
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US-09-454-684-9
US-09-426-571-9
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APPLICANT: Antozak, J.
APPLICANT: Delmedico, M.
APPLICANT: Erickson, J.
APPLICANT: Exickson, J.
APPLICANT: Lambert, D.
APPLICANT: Sista, P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE FUSION-
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
FILE REPERENCE: 7872-073
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                Sequence 25, Application US/09457198A
; GENERAL INFORMATION:
APPLICANT: Christine, Andrews A.
TILE OF INVENTION: Thermostable Enzyme for Nucleic Acid Detection
FILE REFERENCE: Thermostable/nucleic acid detection
CURRENT APPLICATION NUMBER: US/09/457,198A
; CURRENT PELLING DATE: 1999-12-06
; PRIOR PAPLICATION NUMBER: 09/358,972
; RING PAPLICATION NUMBER: 09/358,972
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 25
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CURRENT FILING DATE: 2000-02-29
FRICA APPLICATION NUMBER: 09/315,304
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1589
SOFTWARE: FREUSEQ FOR WINDOWS Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Pyrococcus furiosus
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US-09-515-965-791
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US-09-457-198A-25
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Gaps

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APPLICANT: Kudryk, Bohdan
TITLE OF INVENTION: MONGSPECIFIC ANTIBODY REACTIVE WITH MATRIXMETALLOPROTEINASE CLEAV
TITLE OF INVENTION: PRODUCTS OF FIBRIN (OGEN)
CURRENT APPLICATION NUMBER: US/09/536,998
CURRENT PILING DATE: 2001-04-25
PRIOR APPLICATION NUMBER: 08/900,895
PRIOR FILING DATE: 1997-07-25
NUMBER OF SEQ ID NOS: 3 - 08/900, 895
SOFTWARE: Patentin Version 3.0
SEQ ID NO 1
LENGTH: 5
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; Sequence 9, Application US/09598419
; General INFORMATION:
; APPLICANT: Skeiky, Vasir A.W.
; APPLICANT: Scholler, John
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR TREATMENT AND TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
; FILE REFERENCE: 210121.469C6
; CURRENT APPLICATION NUMBER: US/09/598,419
; CURRENT FILING DATE: 2000-06-20
; NUMBER OF SEQ ID NOS: 357
; SOFTWARE: FastSEQ for Windows Version 3.0/4.0
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APPLICANT: Ropp, Traci M.
APPLICANT: Swanson, Barbara A.
TITLE OF INVENTION: Novel Variant EGIII-Like Cellulase
TITLE OF INVENTION: Compositions
FILE REPERENCE: GC630
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100.0%; Pred. No. 4.2e+06;
ative 0; Mismatches 0;
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100.0%; Pred. No. 4.2e+06;
ative 0; Mismatches 0;
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CURRENT FILING DATE: 2000-08-04
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 5
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US-09-632-426-30
US-09-632-426-30
Sequence 30, Application US/09632426
GENERAL INFORMATION:
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ORGANISM: Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
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US-09-536-998-1
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Best Local Similarity
2; Conserva
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Best Local Similarity
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APPLICANT: Delmedico, M.
APPLICANT: Exickson, J.
APPLICANT: Exickson, J.
APPLICANT: Exickson, J.
APPLICANT: Sista, P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE FUSION-
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
FILE REFERENCE: 7872-073
CURRENT FILING DATE: 2000-02-29
PRIOR PPLICATION NUMBER: 09/915,965A
PRIOR PPLICATION NUMBER: 09/915,304
PRIOR PILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 1994
SOFTHARE: FRAESEQ for Windows Version 3.0
SSQ ID NO 1657
                                                                                                                                                 APPLICANT: SISTA, P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE FUSION-
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
FILE REFERENCE: 7872-073
CURRENT APPLICATION NUMBER: 108/09/515,965A
CURRENT PILING DATE: 2000-02-29
PRIOR APPLICATION NUMBER: 09/315,304
PRIOR APPLICATION NUMBER: 09/315,304
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1994
SOFTWARE: FastSEQ for Windows Version 3.0
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Antczak, J. APPLICANT: Delmedico, M.
                                                                                                                                 Lambert, D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 2; Conserv
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US-09-515-965A-1657
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US-09-515-965A-791
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                                                                                                         Length 5;
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GENERAL INFORMATION:
APPLICANT: Gualfetti, Peter
APPLICANT: Hitchinson, Colin
TITE OF INVENTION: Novel Variant
TITE OF INVENTION: Compositions
TITLE OF INVENTION: Compositions
FILE REFERENCE: GC631
CURRENT APPLICATION NUMBER: US/09/632,570
CURRENT APPLICATION NUMBER: US/09/632,570
CURRENT PILING DATE: 2000-08-04
NUMBER OF SEQ ID NOS: 64
SOFTWARE: FRASESQ for Windows Version 4.0
                                                                                                       DB 20; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 65.2%; Score 15; DB 20; L
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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                                                                                                       65.2%; Score 15; DB ilarity 100.0%; Pred. No. 4.2 Conservative 0; Mismatches
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APPLICANT: WANG, Bryan S.
APPLICANT: PABO, CAZI O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: 105/09/636,243A
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
                 ; LOCATION: (1)...(5)
; OTHER INFORMATION: Xaa = Asn or Gln US-09-632-426-30
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OTHER INFORMATION: Xaa = Asn or Gln
OTHER INFORMATION: BOX1
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ORGANISM: Artificial Sequence
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Best Local Similarity
-hes 2; Conserva
NAME/KEY: VARIANT
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LENGTH: 5
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APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Beter
APPLICANT: Milner, Peter
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karen
APPLICANT: TITLE OF INVENTION: COMPONENTS
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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  0; Mismatches
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CURRENT PAPLICATION NUMBER: US/09/657,276
CURRENT FILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: 60/133,406
PRIOR APPLICATION NUMBER: 60/153,406
PRIOR APPLICATION NUMBER: 60/153,783
PRIOR PILING DATE: 1999-09-10
PRIOR PILING DATE: 1999-09-10
PRIOR PILING DATE: 1999-10-18
NUMBER OF SEC ID NOS: 1617
SEQ ID NO 723
                                                                                                                                                                                                                             APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
TITLE OF INVENTION: DIMERIZING PEPTIDES
TITLE OF INVENTION: DIMERIZING PEPTIDES
CURRENT APPLICATION NUMBER: US/09/642, 234
CURRENT FILING DATE: 2000-08-17
PRIOR PELLING DATE: 1999-08-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 723, Application US/09657276 GENERAL INFORMATION:
                                                                                                                                                                    US-09-642-234-75; Sequence 75, Application US/09642234; GENERAL INFORMATION:
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Best Local Similarity 100...
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SOFTWARE: Patentin Ver. 2.0
  Conservative
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US-09-657-276-723
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                                          4 LW 5
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LENGTH: 5
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  Matches
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65.2%; Score 15; DB 20; Length 5;

Query Match

Best Local Similarity

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RESULT 64
US-09-657-276-1560
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            SOFTWARE: Pat
SEQ ID NO 734
LENGTH: 5
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                                                                                      TYPE: PRT
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APPLICANT: Conjuchem, Inc.
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Alan
APPLICANT: Holmes, Darren
APPLICANT: Holmes, Darren
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karlen
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: COMPONENTS
FILE REFERENCE: 2110
CURRENT APPLICATION NUMBER: 08/09-07
PRIOR FILING DATE: 1999-09-10
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APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Hibaudeau, Raren
APPLICANT: Hibaudeau, Raren
APPLICANT: Hibaudeau, Raren
APPLICANT: Hibaudeau, Resen
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROPTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION NUMBER: US/09/657,276
CURRENT APPLICATION NUMBER: 60/134,406
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR APPLICATION NUMBER: 60/153,406
PRIOR PILING DATE: 1999-05-10
PRIOR PILING DATE: 1999-10-10
PRIOR PILING DATE: 1999-10-18
                                        Gaps
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                                   0; Indels
            Pred. No. 4.2e+06;
100.0%; Preα. ...
-'... 0; Mismatches
                                                                                                                                                                                                                                                       ; Sequence 724, Application US/09657276; GENERAL INFORMATION: APPLICANT: Conjuchem, Inc.
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ORGANISM: Artificial Sequence
                                      2; Conservative
          Best Local Similarity
Matches 2; Conserv
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US-09-657-276-734
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APPLICANT: MORREELS, HENRI
APPLICANT: MORREELS, HENRI
TITLE OF INVENTION: BENZODIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: HESAG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REPERENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Exim. Alan
APPLICANT: Exim. Alan
APPLICANT: Holmes, Daren
APPLICANT: Holmes, Daren
APPLICANT: Holmes, Daren
APPLICANT: Holmes, Daren
APPLICANT: Thibaudeau, Karen
APPLICANT: Thibaudeau, Karen
TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
TITLE OF INVENTION: COMPONERS
FILE REFERENCE: 2110
CURRENT APPLICATION NUMBER: 60/134,406
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR FILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-09-10
PRIOR PRILICATION NUMBER: 60/153,406
PRIOR PRILICATION NUMBER: 60/159,783
PRIOR FILING DATE: 1999-09-10
PRIOR PRILING DATE: 1999-01-0
PRIOR PRILING DATE: 1999-10-18
NUMBER OF SEQ ID NOS: 1617
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                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                     ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: Peptide US-09-657-276-734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: Synthetic; COTHER INFORMATION: Peptide US-09-657-276-1560
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                                                                                                                                                                                                                                                                                                                      Query Match 65.2%; Score 15; DB 20; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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US-09-762-270-19
Sequence 19, Application US/09762270
; GENERAL INFORMATION:
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                                                                                                                           ORGANISM: Artificial Sequence
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Bridon, Dominique
Patentin Ver. 2.1
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APPLICANT: DEFLA, ERIK
APPLICANT: MOEREELS, HERRI
APPLICANT: MOEREELS, HERRI
APPLICANT: MAERTENS, GEERT
TITLE OF INVENTION: HENSOPPRIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
TITLE OF INVENTION: AND USE
FILE REFERENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT PILING DATE: 2001-02-06
PRIOR APPLICATION NUMBER: EP 9897006231
PRIOR APPLICATION NUMBER: EP 98970062.9
PRIOR RILING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: EP 98970062.9
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE PARCHELIN VET. 2.1
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APPLICANT: MOEREELS, HENRI
APPLICANT: MAERTEELS, HENRI
APPLICANT: MAERTEENS, GEERL
TITLE OF INVENTION: BRACODIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: HBSAG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REFERENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT APPLICATION NUMBER: US/09-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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PRIOR APPLICATION NUMBER: PCT/EP99/06231
PRIOR PLLICATION NUMBER: PCT/EP99/06231
PRIOR PLLICATION NUMBER: PP 98870186.8
PRIOR APPLICATION NUMBER: EP 98870186.8
PRIOR APPLICATION NUMBER: EP 99870062.9
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                                                                                                                      ; ORGANISM: Hepatitis B virus US-09-762-270-19
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Sequence 22, Application US/09762270

GENERAL INFORMATION:
APPLICANT: DEPLA, ERIK
APPLICANT: MARENELS, HENRI
APPLICANT: MARENELS, HENRI
TITLE OF INVENTION: BENZODIAZEDINES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: BENZODIAZEDINES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: MAD USE
FILE REFRENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
FILE REPRINGE DATE: 2001-02-06
CURRENT FILING DATE: 1999-08-25
PRIOR FILING DATE: 1999-08-25
PRIOR FILING DATE: 1999-09-01
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PATCHIN UNER: 1999-03-29
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GENERAL INFORMATION:
APPLICANT: DEBLA, ERIK
APPLICANT: MAERIELS, HENRI
APPLICANT: MAERIELS, HENRI
TITLE OF INVENTION: BENZODIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: HBSAG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REFERRANCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT FILING DATE: 2001-02-06
PRIOR APPLICATION NUMBER: PCT/EP99/06231
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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PRIOR APPLICATION NUMBER: PCT/EP99/06231
PRIOR FILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: EP 98870186.8
PRIOR FILING DATE: 1998-09-01
PRIOR PLING DATE: 1999-03-10
PRIOR PLING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PATENTIN Ver. 2.1
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US-09-762-270-22
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US-09-762-270-23
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APPLICANT: Samuel E. Hopkins
APPLICANT: Samuel E. Hopkins
APPLICANT: Samuel E. Hopkins
APPLICANT: Elizabeth M. Dimassimo
APPLICANT: Pamela P. Rusnak
APPLICANT: Thomas Michael Venetta
APPLICANT: NUMBER: 1872-068-99
CURRENT APPLICATION VMMBER: US/09/350,369C
CURRENT FILING DATE: 1999-07-09
NUMBER OF SEQ ID NOS: 2041
                                                                                                                                                                                                                                                                                                               Gaps
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65.2%; Score 15; DB 23; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                          Query Match 65.2%; Score 15; DB 22; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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8.09-350-369C-513
Sequence 513, Application US/09350369C
; GENERAL INFORMATION:
      ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-841-132-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
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US-09-920-306-21
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LENGTH: 5
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Probst, Peter
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121.465C8
CURRENT APPLICATION NUMBER: US/09/841,132
CURRENT FILING DATE: 2001-04-23
NUMBER OF SEQ ID NOS: 599
SOFTWARE: PastSEQ for Windows Version 3.0/4.0
SEQ ID NO 9
LENGTH: 5
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APPLICANT: Guthrie, Kelly
APPLICANT: Guthrie, Kelly
APPLICANT: Merutka, Mohmed
APPLICANT: Anwer, Mohmed
APPLICANT: Lambert, Dennis
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES
TITLE REPERRICE: 7872-043
CURRENT APPLICATION NUMBER: 05/09/834,784
CURRENT FILING DATE: 2001-04-13
PRIOR PILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
SOFTWARE: FREUSED for Windows Version 3.0
PRIOR FILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: EP 98870186.8
PRIOR FILING DATE: 1999-09-01
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFWARE: Patentin Ver. 2.1
SEQ ID NO 23
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; OTHER INFORMATION: Core polypeptide
US-09-834-784-791
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; Sequence 791, Application US/09834784
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 2; Conservat
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APPLICANT: Samuel E. Hopkins
APPLICANT: Samuel E. Hopkins
APPLICANT: Balizabeth M. Dimassimo
APPLICANT: Tolar Samuel P. Rusnab
APPLICANT: Thomas Michael Venetta
APPLICANT: Thomas Michael Venetta
TITLE OF INVENTION: Methods and Compositions for Detection of HIV GP41 Peptides
FILE REFERENCE: 7872-068-999
CURRENT APPLICATION NUMBER: US/09/350,369C
CURRENT FILING DATE: 1999-07-09
NUMBER OF SEQ ID NOS: 2041
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 5
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Sequence 36, Application US/10005305

GENERAL INFORMATION:
APPLICANT: The Government of the United States of America, represented by the APPLICANT: Secretary, Department of Health and Human Services
TITLE OF INVENTION: T20/DP178 AND T21/DP107 ARE ACTIVATORS
TITLE OF INVENTION: OF HUMAN PHAGOCYTE
TITLE OF INVENTION: PORMYL PEPTIDE RECEPTORS
FILE REFERENCE: NH171.001C1 US/10/005,305
CURRENT APPLICATION NUMBER: US/10/005,305
PRIOR PILING DATE: 2000-05-05
PRIOR FILING DATE: 2000-06-05
PRIOR FILING DATE: 1999-05-05
NUMBER OF SEQ ID NOS: 203
SEQ ID NO 36
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 65.2%; Score 15; DB 23; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Human Immunodeficiency Virus-1 (HIV-1)
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                                                                                                                        Sequence 1389, Application US/09350369C GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: Artificial Peptide US-10-005-305-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                  US-09-350-369C-1389
1 LW 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
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                                                                        RESULT 74
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398, App
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211, App
217, App
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(without alignments)
23.274 Million cell updates/sec
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score greater than or equal to the score of the result beir
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                         GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-636-243B-75
US-09-968-744A-211
US-09-968-744A-211
US-09-968-744A-213
US-10-174-717A-94
US-10-099-056-2683
US-10-099-056-2683
US-10-05-33556-34
PCT-US02-33556-34
PCT-US02-33556-34
PCT-US02-33556-36
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PCT-US02-34760-47
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PCT-US02-34760-47
PCT-US02-34760-47
PCT-US02-24469A-21
PCT-US02-24469A-21
PCT-US02-24469A-21
PCT-US02-24310-166
PCT-US02-33507-23
PCT-US02-33507-23
PCT-US02-3410-15
                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                            425801 segs, 83787487 residues
                                                                                                                                                                                                                                                                                                                                                                              SUMMARIES
                                                                                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries
                                                           February 12, 2003, 10:51:59
                                           - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
                                                                                               US-09-403-440A-1
23
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Match Length
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                                                                                                                                  Scoring table:
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Perfect score:
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No.
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116, Appl 1145, Appl 1152, Appl 1152, Appl 1152, Appl 118, Appl 118, Appl 119, Appl 110, Appl 110, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 117, Appl 118, Appl 119, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 117, Appl 118, Appl 119, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 117, Appl 118, Appl 119, Appl 119, Appl 111, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 117, Appl 118, Appl 119, Appl 119, Appl 111, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 117, Appl 117, Appl 118, Appl 119, Appl 119, Appl 119, Appl 111, Appl 111, Appl 111, Appl 111, Appl 112, Appl 113, Appl 114, Appl 115, Appl 116, Appl 117, Appl 11

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Sequence 211, Application US/09968744A

Sequence 211, Application US/09968744A

Sequence 211, Application US/09968744A

Sequence 211, Application US/09968744A

APPLICANT: Tominson, Ian M

APPLICANT: Minter, Gregory

TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands

FILE REFERENCE: 8039/1073

CURRENT APPLICATION NUMBER: US/09/968,744A

CURRENT FILING DATE: 2003-01-13

PRIOR APPLICATION NUMBER: US 60/065,248

PRIOR APPLICATION NUMBER: US 60/066,729

PRIOR PILING DATE: 1997-11-21

PRIOR PLING DATE: 1998-10-20

PRIOR PLING DATE: 1998-10-20

PRIOR PLING DATE: 1998-10-20

PRIOR FILING DATE: 1998-10-20

SPRIOR FILING DATE: 1998-10-20
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**REPERCENCE:

**Sequence:

**APPLICANT: Tomlinson, Tan M

**APPLICANT: Tomlinson, Tan M

**APPLICANT: Tomlinson, Tan M

**APPLICANT: Winter, Gregory

**TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands

**TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands

**TITLE OF INVENTION: Member: US/09/968,744A

**CURRENT APPLICATION NUMBER: US 9722131.1

**PRIOR FILING DATE: 1997-10-20

**PRIOR APPLICATION NUMBER: US 60/065,248

**PRIOR APPLICATION NUMBER: US 60/066,729

**PRIOR APPLICATION NUMBER: US 60/066,729

**PRIOR PILING DATE: 1997-11-21

**PRIOR PILING DATE: 1998-10-20

**PRIOR FILING DATE: 2000-02-4

**NUMBER OF SEQ ID NOS: 350

**PRIOR FILING DATE: 2000-03-14

**PRIOR FILING DATE: 2000-03-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
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US-09-968-744A-217
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US-09-968-744A-217
                                                                                                                                                                     RESULT 3
US-09-968-744A-211
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       4 LW 5
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APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Fetterick, Robert
APPLICANT: Fetterick, Robert
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, National
APPLICANTION: COACTIVAOR BINDING
FILE REFERENCE: 9811-008-999
CURRENT APPLICATION NUMBER: US/09/281,717A
FILE REPERENCE: 1999-03-30
FRIOR APPLICATION NUMBER: US 60/079,956
PRIOR PRILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3
LENGTH: 5
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   Sequence 38, Appl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 82.6%; Score 19; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 5; Conservative 0; Mismatches 0; Indels
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6 US-10-058-513-38
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FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SEQ ID NO 75
                                                                                                    ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (2)...(3)
; OTHER INFORMATION: Xaa is any amino acid US-09-281-717A-3
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                                                                                                                                                                                                                                                               Sequence 3, Application US/09281717A GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                    APPLICANT: Baxter, John
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 2; Conserva
47.8
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PRIOR FILING DATE: 2001/03/14
NUMBER OF SEQ ID NOS: 2850
SOFTWARE: PatentIn version 3.1
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MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                   Query Match
Best Local Similarity 50.v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                                                                  TYPE: PRT
ORGANISM: Homo Sapiens
FEATURE:
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US-10-174-717A-94
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                                                          SEQ ID NO 2683
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TITLE OF INVENTION: More and more protein-protein interactions in adipocyte cells
TITLE REFERENCE: B4838A
CURRENT APPLICATION NUMBER: US/10/099,056
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: US 60/275.734
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0
                                                                                                                                                                                                                                                         ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, St. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 56.5%; Score 13; DB 6; Length 5; Best Local Similarity 50.0%; Pred. No. 3.5e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                         Sequence 51, Application US/10174717A
APPLICANT: Koide, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 18-Jun-2002
PRIOR APPLICATION DATA:
FILING DATE: 18-Jun-2002
APPLICATION NUMBER: 09/096,749
FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY/AGRY INFORMATION:
NAME: Ann S. Vikshins
REGISTRATION NUMBER: 37,748
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORIGINAL SOURCE:
SEQUENCE DESCRIPTION: SEQ ID NO: 51:
US-10-174-717A-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2683, Application US/10099056
GENERAL INFORMATION:
APPLICANT: Hybrigenics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (612) 339-3
INFORMATION FOR SEQ ID NO: 51:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                              ZIP: 55402
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                        STATE: MIN
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Sequence 94, Application US/10174717A

TITLE OF INVENTION: ARTHFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118

CORRESPONDENCE ADDRESS:
ADDRESSE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, St. 1600

CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 52.2%; Score 12; DB 6; Length 5; Best Local Similarity 50.0%; Pred. No. 3.5e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                      Score 13; DB 6; Length 5;
Pred. No. 3.5e+05;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY AGENT INFORMATION:
NAME: Ann S. Vikenins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
; OTHER INFORMATION: Translation of SEQ ID NO:691
US-10-099-056-2683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
OPREATING SYSTEM: WINDOWS
SOFTWARE: FaatSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/174,717A
PILING DATE: 18-Jun-2002
PRIOR APPLICATION NUMBER: 09/096,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT TYPE: internal ORIGINAL SOURCE: SEQUENCE DESCRIPTION: SEQ ID NO: 94:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (612) 373-690
TELEFAX: (612) 339-3061
INFORMATION FOR SEQ ID NO: 94:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
                                                                      56.5%;
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APPLICANT: Amgen, Inc.
TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferor-Gamma
TITLE OF INVENTION: Neutralizing Activity
FILE REFERENCE: A-799
CURRENT APPLICATION NUMBER: PCT/US02/33556
CURRENT FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 135
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Amgen, Inc.
TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma TITLE OF INVENTION: Neutralizing Activity
FILE REFERENCE: A-799
CURRENT APPLICATION NUMBER: PCT/USO2/33556
CURRENT FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 135
SOFTWARE: Patentin version 3.0
SEQ ID NO 36
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Amgen, Inc.
TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma TITLE OF INVENTION: Neutralizing Activity
FILE REPERENCE: A-799
CURRENT APPLICATION NUMBER: PCT/US02/33556
CURRENT FILIG DARE: 2002-10-17
NUMBER OF SEQ 1D NOS: 135
SOFTWARE: PatentIn version 3.0
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Query Match

47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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  PCT-US02-33556-34; Sequence 34, Application PC/TUS0233556; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              // Sequence 36, Application PC/TUS0233556
// GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 100.0
Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; ORGANISM: Homo sapiens
PCT-US02-33556-36
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; ORGANISM: Homo sapiens
PCT-US02-33556-40
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Best Local Similarity
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                                                                                                                                                                                                                                             APPLICANT: Houghton, Alan
APPLICANT: Oustreelli, Ouathek
APPLICANT: Oustreelli, Ouathek
TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
CURRENT APPLICATION NUMBER: US/10/053,498B
CURRENT FILING DATE: 2002-01-17
PRIOR FILING DATE: 1997-10-31
NUMBER OF SEQ ID NOS: 321
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 320
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: AMGEN INC.

TITLE OF INVENTION: SPECIFIC BINDING AGENTS OF HUMAN ANGIOPOIETIN-2
FILE REFERENCE: A-801B
CURRENT APPLICATION NUMBER: PCT/USO2/32657
CURRENT FILING DATE: 2002-10-11
PRIOR APPLICATION NUMBER: US 60/414,155
PRIOR PILING DATE: 2002-09-27
PRIOR PILING DATE: 2001-10-11
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 359
SOFTWARE: Patentin version 3.1
SEQ ID NO 65
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
52.2%; Score 12; DB 6; Length 5;
Best Local Similarity 40.0%; Pred. No. 3.5e+05;
Matches 2; Conservative 0; Mismatches 3; Indels
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                            APPLICANT: Sloan-Kettering Institute for Cancer Research APPLICANT: Rothman, James E. APPLICANT: Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: Random peptide in coliphage M13
US-10-053-498B-320
                                                                                                                           ; Sequence 320, Application US/10053498B ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 65, Application PC/TUS0232657; GENERAL INFORMATION:
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                                                                                                                                                                                                       Rothman, James E.
Mayhew, Mark
                                                                                                      US-10-053-498B-320
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SEGRENEAL INFORMATION:
APPLICANT: Abbott Laboratories
APPLICANT: Abbott Laboratories
APPLICANT: Bradley, Michael F.
TITLE OF INVENTION: TETRA-, PENTA-, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: HAVING ANTIANGIOGENIC ACTIVITY
FILE REFERENCE: 6854.00.01
CURRENT APPLICATION NUMBER: PCT/US02/34760
CURRENT PLING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: 10/263,811
PRIOR PILING DATE: 2002-10-04
PRIOR PELLING DATE: 2001-10-14
PRIOR PELLING DATE: 2001-10-14
PRIOR FILING DATE: 2001-10-31
PRIOR APPLICATION NUMBER: 10/000,007
PRIOR FILING DATE: 2001-10-31
PRIOR PELLING DATE: 2001-10-31
SROFTWARE: FASELESQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: AATA'S, COTTUDA
APPLICANT: Bradley, Michael F.
TITLE OF INVENTION: TETRA, PENTA, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: TETRA, PENTA, PENTA,
                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 5;
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                                                              Length 5;
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CTHER INFORMATION: Xaa = proethylamide at position 5
PCT-US02-34760-46
                                                          47.8%; Score 11; DB 1; Le 100.0%; Pred. No. 3.5e+05; ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
47.8%; Score 11; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
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OTHER INFORMATION: Antiangiogenic Peptide
                                                                                                                                                                                                                                                                                                                                                                                          PCT-1802-34760-46; Sequence 46, Application PC/TUS0234760; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                Conservative
                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: VARIANT
PCT-US02-34760-21
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; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
PCT-US02-33985-2
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APPLICANT: Bradley, Michael F.
ITILE OF INVENTION: FEFRA-, PENTA-, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: FFFRA-, PENTA-, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: HAVING ANTIANGIOGENIC ACTIVITY
FILE REFERENCE: 6854.WO.01
CURRENT APPLICATION NUMBER: PCT/US02/34760
CURRENT PILING DATE: 2002-10-30
PRIOR FILING DATE: 2002-10-04
PRIOR FILING DATE: 2001-10-31
PRIOR PELICATION NUMBER: 10/000,007
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
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   0; Indels
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OTHER INFORMATION: Xaa = prolylethylamide at position 5
                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: 16 Human Secreted Proteins
FILE REFERENCE: PS739PCT
   0; Mismatches
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OTHER INFORMATION: Xaa = Nva at position 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: PCT/US02/33985
CURRENT FILING DATE: 2002-10-24
PRIOR APPLICATION NUMBER: 60/330,629
PRIOR FILING DATE: 2001-10-26
NUMBER OF SEQ ID NOS: 62
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO. 2
LENGTH: 5
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                                                                                                                                                                                                                                                                                               Sequence 2, Application PC/TUS0233985 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Abbott Laboratories
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ORGANISM: Artificial Sequence
   Conservative
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATING TO THE DAPTOMYCIN
TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATING TO THE DAPTOMYCIN
TITLE OF INVENTION: BIOSYNTHETIC GENE CLUSTER
FILE REFERENCE: CUB-1.2 PCT CIP
CURRENT APPLICATION NUMBER: PCT/US02/24310
CURRENT APLICATION NUMBER: PCT/US01/32354
PRIOR APPLICATION NUMBER: PCT/US01/32354
PRIOR APPLICATION NUMBER: 60/310,385
PRIOR APPLICATION NUMBER: 60/310,385
PRIOR PILING DATE: 2001-08-06
PRIOR PILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 170
SOFTWARE: PATENTIN DATE: 202-05-10
                                                                                                              0; Gaps
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                                                        Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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Query Match

47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                              FUEL SEGUENCE 21, Application PC/TUS0224469
; GENERAL INFORMATION:
; APPLICANT: METCK & Co., Inc.
; TITLE OF INVENTION: Biological Assay Detection Method
; FULE REFERENCE: PCT 20900'
; CURRENT APPLICATION NUMBER: PCT/US02/24469
; CURRENT FILING DATE: 2002-08-02
; PRIOR PELING DATE: 2001-08-07
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: FastSEQ for Windows Version 4.0
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OTHER INFORMATION: Xaa = Any Amino Acid
FEATURE:
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; OTHER INFORMATION: Xaa = Any Amino Acid
PCT-US02-24469-21
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APPLICANT: CUBIST PHARMACEUTICALS, INC.
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ORGANISM: Artificial Seguence
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NAME/KEY: VARIANT
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     PCT-US02-16525A-26
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LENGTH: 5
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Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REFERENCE: PFS.56PT.
CURRENT APPLICATION NUMBER: PCT/US02/16525A
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR FILING DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
LENGTH: 5
                                                                                                                                                   NAME/KEY: VARIANT
LOCATION: (5)...(5)
CTHER INFORMATION: Xaa = proethylamide at position 5
PCT-US02-34760-47
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SERERAL INFORMATION:
SETTLE APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: 41 Human Secreted Proteins
FILE REFERENCE: PS740FCT
CURRENT APPLICATION NUMBER: PCT/US02/35606
CURRENT FILING DATE: 2002-11-06
PRIOR APPLICATION NUMBER: 60/331,046
PRIOR PILING DATE: 2001-11-07
NUMBER: PATENT FILING DATE: 2001-11-07
NUMBER: PATENT FILING DATE: 2001-11-07
NUMBER: PATENT FILING DATE: 2001-11-07
                                                                                                        OTHER INFORMATION: Antiangiogenic Peptide
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; GENERAL INFORMATION:
                         TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Homo Sapiens
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Matches 1; Conserva
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LOCATION: (3)
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PCT-US02-35606-2
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LENGTH: 5
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LENGTH: 5
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                                                                            FEATURE:
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Sequence 23, Application PC/TUS0232307
GENERAL INFORMATION:
APPLICANT: IMMUNOMEDICS, INC.
TITLE OF INVENTION: THERAPETIC AGENTS
TITLE OF INVENTION: THERAPETIC AGENTS
TITLE OF INVENTION: THERAPETIC AGENTS
FILE REFERENCE: 018733-1148
CURRENT APPLICATION NUMBER: PCT/US02/32307
CURRENT APPLICATION NUMBER: PCT/US02/32307
CURRENT APPLICATION NUMBER: US 60/416,531
PRIOR FILING DATE: 2002-10-08
NUMBER OF SEQ ID NOS: 27
SOGTWARE: Patentin version 3.1
SEQ ID NO 23
LENGTH: 5
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Sequence 5, Application PC/TUS0239643
GENERAL INFORMATION:
APPLICANT: Smith, Thomas M
APPLICANT: Kirley, Terence L
TITLE OF INVENTION: APPRASE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING SAME
FILE REFERENCE: 0091830/0502531
CURRENT APPLICATION NUMBER: PCT/US02/39643
CURRENT PILING DATE: 2002-12-12
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.1
SEQ ID NO 5
LENGTH: 5
          ; OTHER INFORMATION: Description of Artificial Sequence:/note; OTHER INFORMATION: Synthetic Construct
PCT-US02-39109-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 47.8%; Score 11; DB 1; Length 5; Local Similarity 100.0%; Pred. No. 3.5e+05; nes 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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PCT-US02-40891-1113
F Sequence 1113, Application PC/TUS0240891
GENERAL INPORMATION:
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ORGANISM: Murinae gen. sp.
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ORGANISM: Homo sapiens
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CURRENT PAPLICATION NUMBER: PCT/US02/39109
CURRENT FILING DATE: 2002-12-06
PRIOR APPLICATION NUMBER: 60/338,971
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; OTHER INFORMATION: Description of Artificial Sequence: Illustrative; OTHER INFORMATION: thioesterase motif PCT-US02-24310-166
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                                                                                                             Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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ITLEO PO INVENTION: Biological Assay Detection Method
FILE REFERENCE: PCT 20000Y
CURRENT APPLICATION NUMBER: PCT/US02/24469A
CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: 60/310,599
PROR FILING DATE: 2001-08-07
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 15
LENGTH: 5
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OTHER INFORMATION: Xaa = Any Amino Acid
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COCATION: (1)...(5)
OTHER INFORMATION: Xaa = Any Amino Acid
PCT-US02-24469A-21
                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 21, Application PC/TUS0224469A; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
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; Sequence 14, Application PC/TUS0300440; GENERAL INFORMATION:
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                    PRIOR FILING DATE: 2002-11-05
                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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Best Local Similarity
Thes 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 1; Conserv
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                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCT-US02-40892-398
                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (3)
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                                                                                                                                                                                                         SEQ ID NO 398
LENGTH: 5
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LENGTH: 5
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OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
APPLICANT: Human Genome Sciences, Inc.
TILE OF INVENTION: Albumin Fusion Proteins
FILE REPERENCE: PF5.4Pc7
CURRENT APPLICATION NUMBER: PC7/US02/40891
CURRENT FILING DATE: 2002-12-23
FRIOR PLING DATE: 2002-02-8
FRIOR APPLICATION NUMBER: 60/360,000
FRIOR FILING DATE: 2002-02-28
FRIOR PELING DATE: 2002-05-10
FRIOR PELING DATE: 2002-10-10
FRIOR PELING DATE: 2002-10-10
FRIOR PELING DATE: 2002-10-10
FRIOR PELING DATE: 2002-10-11
FRIOR PELING DATE: 2002-10-11
FRIOR PELING DATE: 2002-10-11
FRIOR PELING DATE: 2002-10-11
FRIOR APPLICATION NUMBER: 60/423,623
FRIOR PELING DATE: 2002-10-28
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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FILE REFERENCE: PF548CT
CURRENT APPLICATION NUMBER: PC7/US02/40892
CURRENT FILING DATE: 2002-12-23
PRIOR APPLICATION NUMBER: 60/341,811
PRIOR APPLICATION NUMBER: 60/340,
PRIOR APPLICATION NUMBER: 60/378,950
PRIOR APPLICATION NUMBER: 60/378,950
PRIOR PILING DATE: 2002-02-18
PRIOR PILING DATE: 2002-05-10
PRIOR PILING DATE: 2002-07-44
PRIOR FILING DATE: 2002-07-44
PRIOR FILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-10-18
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PRIOR APPLICATION WINBER: 60/420,246
PRIOR FILING DATE: 2002-10-23
PRIOR APPLICATION NUMBER: 60/423,623
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LENGTH: 5
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OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
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PRIOR APPLICATION NUMBER: 60/350,358
PRIOR FILING DATE: 2002-01-24
Remaining Prior Application data removed - See File Wrapper or PALM.
WUMBER OF SEQ ID NOS: 858
SOFTWARE: Patentin Ver. 2.0
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APPLICANT: Sagres Discovery
APPLICANT: Engelhard, Eric K.
APPLICANT: Engelhard, Eric K.
APPLICANT: Engelhard, Eric K.
APPLICANT: Morris, David W.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER FILE REFERENCE: 52945200014
CURRENT APPLICATION NUMBER: PCT/US02/33835
PRIOR FILING DATE: 2002-12-19
PRIOR FILING DATE: 2001-10-23
NUMBER OF SEC 1D NOS: 61
SOFTWARE: PatentIn version 3.1
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OTHER INFORMATION: "Xaa" at position 3 can be any amino acid.
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TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30
FILE REFERENCE: MXI-190PC
CURRENT APPLICATION NUMBER: PCT/US03/00440
PRIOR APPLICATION NUMBER: US 60/347649
PRIOR PILING DATE: 2002-01-09
PRIOR PILING DATE: 2002-01-09
PRIOR PLILNG DATE: 2002-01-09
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APPLICANT: JONCZYK, ALERED
APPLICANT: GOCDMAN, SIMON
APPLICANT: GCODMAN, SIMON
APPLICANT: DIEFENBACH, BEATE
APPLICANT: KESSLER, HORST
APPLICANT: FINSINGER, DIRK
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR PILICATION NUMBER: 08/710,251
PRIOR FILING DATE: 1996-09-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/627,173
FILING DATE: 03-APR-1996
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 2201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS Word
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-08-832-443C-13
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          PCT-US03-00440-38
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GENERAL INFORMATION:

APPLICANT: Medarex, Inc. et al.

TILLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30
FILE REPERENCE: MXI-180PC

CURRENT APPLICATION NUMBER: PCT/US03/00440

CURRENT PILING DATE: 2003-01-07

PRIOR PILING DATE: 2002-01-09

PRIOR PILING DATE: 2002-01-09

PRIOR FILING DATE: 2002-08-19

NUMBER OF SEQ ID NOS: 53

SOFTWARE: FASTERQ for Windows Version 4.0

SEQ ID NO 26
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TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30
FILE REFERENCE: MXI-180PC
CURRENT APPLICATION NUMBER: PCT/US03/00440
CURRENT FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: US 60/347649
PRIOR PELING DATE: 2002-01-09
PRIOR FILING DATE: 2002-01-09
PRIOR FILING DATE: 2002-01-09
PRIOR FILING DATE: 2002-10-09
PRIOR FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FASELSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                          Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 2002-08-19
PRIOR APPLICATION NUMBER: US 60/431684
PRIOR FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 53
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 14
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                                                                                                                                                   LENGTH: 5
; TYPE: PRT
; ORGANISM: Homo sapiens
PCT-US03-00440-14
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; TYPE: PRT
; ORGANISM: Homo sapiens
PCT-US03-00440-26
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ORGANISM: Homo sapiens
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US-08-812-443C-13
US-08-812-443C-13
SEQUENCE 13, Application US/08832443C
GENERAL INFORMATION:
APPLICANT: TSYRLOVA, IRENA
NOLPE, STEPHEN D
TITLE OF INVENTION: INHIBITOR OF STEM CELL PROLIFERATION AND
USES THEREOF
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Query Match
47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
REPLICATION UNMERS: US/08/832,443C
FILING DATE: 03-Apr-1997
CLASSIFICATION: «Unknown»
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                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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GENERAL INFORMATION:
APPLICANT: GOODAAN, SIMON
APPLICANT: GOODAAN, SIMON
APPLICANT: GOODAAN, SIMON
APPLICANT: ELSEER, HORST:
APPLICANT: FINSINGER, HORST:
APPLICANT: FINSINGER, DIRK
ITILE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-185.1
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
FRIOR FILING DATE: 1997-08-07
FRIOR FILING DATE: 1996-09-13
NUMBER OF SEQ ID NOS: 175
SEQ ID NO 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: JONCZYK, ALFRED
APPLICANT: GONDANA, SIMON
APPLICANT: GONDANA, SIMON
APPLICANT: BIEFENBACH, BEBATE
APPLICANT: KESSLER, HORST
APPLICANT: KESSLER, HORST
APPLICANT: FINSINGER; DIRK
TILLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATENTIN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 152, Application US/08908416B; GENERAL INFORMATION:
                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 147
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US-08-908-416B-157
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                                                                        OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                     Query Match 47.8%; Score 11; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 162, Application US/08908416B
Sequence 162, Application US/08908416B
GENERAL INFORMATION:
APPLICANT: JONCZYK, ALFRED
APPLICANT: DISCHENBACH, BEATE
APPLICANT: DISCHENBACH, BEATE
APPLICANT: FISSINGEN, HORST
APPLICANT: FINSINGEN, DIRK
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT APPLICATION NUMBER: US/08/016
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
SOFTWARE: PatentIn Ver: 2.1
SEQ ID NO 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: GOODMAN, SIMON
APPLICANT: GOODMAN, SIMON
APPLICANT: DIEFNBACH, BEATE
APPLICANT: KESSLER, HORST
APPLICANT: FINSINGER, DIRK
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REPERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION DATE: 1996-09-13
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATENTIN OS: 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 167, Application US/08908416B; GENERAL INFORMATION:
APPLICANT: JONCZYK, ALFRED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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US-08-908-416B-167
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LENGTH: 5
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APPLICANT: Fettig, Wolfgang J.
APPLICANT: Lenter, Martin U.
APPLICANT: Deter, Martin U.
APPLICANT: Garin-Chesa, Pilar
APPLICANT: Garin-Chesa, Pilar
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, John E.
APPLICANT: Telan, Leila A.
TITLE OF INVENTION: FAP-Activated Anti-Tumour Compounds;
FILE REFERENCE: 1/1097 a.
CURRENT APPLICATION NUMBER: US/09/533,800
CURRENT APPLICATION A.
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     SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/912,951
FILING DATE: 14.AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14.AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01.OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01.OCT-1997
                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1997
PRILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
                                                                                                                                                                                                                                                                          FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
FILING DATE: 06-MAX-1997
FILING DATE: 06-MAX-1997
APPLICATION NUMBER: US 08/851,843
APPLICATION NUMBER: US 08/854,050
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Matches 1; Conservative 0; Mismatches
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APPLICATION NUMBER: US 08/911,312
                            CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/721,456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 540:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: US 60/134,136
                                                                           FILING DATE: 22-Nov-2000 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 16, Application US/09533800
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 14-AUG-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
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US-09-533-800-16
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FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
OTHER INFORMATION: peptide
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Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                          Query Match

47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: JONGEXY, ALFRED
APPLICANT: GOODWAN, SIMON
APPLICANT: GOODWAN, HORST
APPLICANT: BISSLER, HORST
APPLICANT: FINSINGER, DIRK
ITILE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851;
APPLICANT: FINSINGER, DIRK
ITILE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851;
AURRENT APPLICATION NUMBER: US/08/908,416B;
CURRENT APPLICATION NUMBER: 08/710,251
PRIOR FILING DATE: 1996-09-13
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATENTIN VET. 2.1
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 38
US-09-721-456-540
US-09-721-456-540, Application US/09721456
GENERAL INFORMATION:
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Morin, Gregg B.
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STATE: California
COUNTRY: USA
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ORGANISM: Artificial Sequence
                                                                         US-08-908-416B-167
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US-08-908-416B-172
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LENGTH: 5
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GENERAL INFORMATION:

APPLICANT: RAZAZIAN, HAIG H.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOENE, JEF D.

APPLICANT: BETH, DOMBROSKI A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF MAMMALIAN RETROTRANSPOSONS

FILE REFERENCE: 9596-23U3 (053893-5006-02)

CURRENT APPLICATION NUMBER: US(99/653,8128)

CURRENT FILING DATE: 1200-09-01

PRIOR PILING DATE: 1297-04-28

PRIOR PILING DATE: 1996-11-15

PRIOR FILING DATE: 1996-11-16

NUMBER OF SEQ ID NOS: 152

SOFTWARE: PATENTIN VERSION 3.1

SEQ ID NO 145

LENGTH: S
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APPLICANT: KAZAZIAN, HAIG H.
APPLICANT: KAZAZIAN, HAIG H.
APPLICANT: MORAN, JOHN V.
APPLICANT: MORAN, JOHN V.
APPLICANT: BETH, DOMENCSKI A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF MAMMALIAN RETROTRANSPOSONS
FILE REPERBENCE: 9596-2313 (053893-5006-02)
CURRENT APPLICATION NUMBER: US/09/653,812B
CURRENT FILING DATE: 2000-09-01
PRIOR PLILNG DATE: 1997-04-28
PRIOR PLILNG DATE: 1997-04-28
PRIOR PLILNG DATE: 1996-11-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Wild type L1.2mneoI construct
                                                                                                                                                             LOCATION: (3)..(3); OTHER INFORMATION: Xaa equals any amino acid US-09-726-348B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 145, Application US/09653812B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          // Sequence 152, Application US/09653812B
// GENERAL INFORMATION:
                                                                                        OTHER INFORMATION: WSXWS motif
FEATURE:
NAME/KEY: SITE
                        TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-653-812B-145
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                                                                     FEATURE:
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TITLE OF INVENTION: Novel Methods Of Ultrasound Treatment Using Gas Or Gaseous Precur
TITLE OF INVENTION: Filled Compositions
FILE REFERENCE: UNGRISCO CURRENT APPLICATION NUMBER: US/09/813,484
CURRENT FILING DATE: 2001-03-21
PRIOR FILING DATE: 1997-09-15
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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GENERAL INFORMATION:
APPLICANT: Wei, Ying-Fei
TITLE OF INVENTION: Transforming Growth Factor Alpha HIII
TITLE OF INVENTION UNBER: US/09/726,348B
CURRENT APPLICATION NUMBER: US/09/726,348B
CURRENT PILING DATE: 1990-12-01
PRIOR APPLICATION NUMBER: 08/778,545
PRIOR APPLICATION NUMBER: 06/011,136
PRIOR FILING DATE: 1999-01-03
PRIOR FILING DATE: 1999-10-04
PRIOR FILING DATE: 1999-12-02
NUMBER: OF SEQ ID NOS: 21
SOFTWARE: Patentin Version 3.0
SOFTWARE: Patentin Version 3.0
                                                                                                                                                                                                                                47.8%; Score 11; DB 5; Length 5; ilarity 100.0%; Pred. No. 3.5e+05; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: misc_feature

LOCATION: (4)...(4)

COTHER INFORMATION: Xaa is azetidine

US-09-813-484-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-813-484-1; Sequence 1, Application US/09813484; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 28
SOFTWARE: Patentin version 3.1
                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                             OTHER INFORMATION: Precursor
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Best Local Similarity
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US-09-726-348B-5
                                                                                                                                                                                 US-09-533-800-16
                                              SEQ ID NO 16
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                                                                                             TYPE: PRT
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                                                                                                                                     FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: MEDLEN & CARROLL, LLP
STREET: 220 Montgomery Street, Suite 2200
CITY: San Francisco
STATE: CA
               ;
0
                                                                                                                                                                                                                                  FILLE OF INVENTION: Human Frezzled-Like Protein FILLE REFERENCE: PF380C1
CURRENT APPLICATION NUMBER: US/09/610,650B
CURRENT FILING DATE: 2000-06-30
FRIOR APPLICATION NUMBER: 09/132,315
FRIOR FILING DATE: 1998-08-11
FRIOR FILING DATE: 1997-08-12
FRIOR FILING DATE: 1998-04-10
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.1
SOFTWARE: PatentIn version 3.1
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100.0%; Pred. No. 3.5e+05;
tive 0; Mismatches 0;
             Mismatches
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CLASSIFICATION: «UNKNOWN»
PRIOR APPLICATION NUMBER: US/09/307,579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/943,120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COCATION: (3)...(3)
COTHER INFORMATION: Xaa is any amino acid
US-09-610-650B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 07-May-1999
APPLICATION NUMBER: 08/884,325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/09943120; GENERAL INFORMATION:
APPLICANT: Rothschild, Kenneth J.
                                                                                                                                                                                               ; Sequence 18, Application US/09610650B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
             0;
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ZIP: 94104
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: MISC_FEATURE
                                                                                                                                                                                US-09-610-650B-18
             1;
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           Matches
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COMPUTER: I FALPLY GLEAR
COMPUTER: I BM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT PAPLICATION DATA:
APPLICATION NATE: US/09/556,972
FILING DATE: 24-Apr-2000
PRIOR APPLICATION NUMBER: US/08/996,338
FILING DATE: 22-DEC-1997
APPLICATION NUMBER: UP 215,488/1997
FILING DATE: 28-UU-1997
APPLICATION NUMBER: UP 215,488/1997
FILING DATE: 28-UU-1997
APPLICATION NUMBER: UP 291,837/1997
FILING DATE: 09-OCT-1997
APPLICATION NUMBER: UP 291,837/1997
FILING DATE: 09-OCT-1997
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                        47.8%; Score 11; DB 5; Length 5; 100.0%; Pred. No. 3.5e+05; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 419 Seventh Street, N.W., Suite 300 CITY: Washington
                                                                                                                                                                                           ; OTHER INFORMATION: Mutant L1.2mmeoI construct US-09-653-812B-152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
TOPOLOGY: linear
MOLECULE TYPE: peptide
FRAGMENT TYPE: internal fragment
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
US-09-556-972-12
                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 44
US-09-556-972-12
; Sequence 12. Application US/09556972
; GENERAL INFORMATION:
APPLICANT: TORIGOE, Kakuji
CKURA, Takanori
HITLE OP INVENTION: POLYPEPTIDES
INUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: BROWDY AND NEIMARK
PRIOR APPLICATION NUMBER: US 60/006,831
PRIOR FILING DATE: 1955-11-16
NUMBER OF SEQ ID NOS: 152
SOFTWARE: Patentin version 3.1
SEQ ID NO 152
LENGTH: 5
TYPE: PRT
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REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: TOR
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                                                                                                                                                   ORGANISM: Artificial Sequence
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SEQUENCE CHARACTERISTICS
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Matches 1; Conservative
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COUNTRY: USA
ZIP: 20004
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; Sequence 20, Application US/09969748C; GENERAL INFORMATION:
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TITLE OF INVENTION: URGS THEROY.

TITLE OF INVENTION: USES THEREOF.

TITLE OF INVENTION: USES THEREOF.

FILE REFERENCE: 076518-0150

CURRENT APPLICATION NUMBER: 02/09/936,759

CURRENT APPLICATION NUMBER: PCT/USO/07107

PRIOR PILING DATE: 2000-03-16

PRIOR PILING DATE: 1999-03-17

PRIOR PREDICATION NUMBER: 09/149,927

PRIOR PILING DATE: 1999-09

PRIOR PILING DATE: 1999-09

PRIOR PILING DATE: 1999-09

PRIOR PILING DATE: 1997-09-09

NUMBER OF SEQ ID NOS: 120

SEQ ID NOS: 120
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                         NAME: CATTOIL, PETER G.
REGISTRATION NUMBER: 32,837
REGISTRATION NUMBER: 32,837
RELECOMMUNICATION INFORMATION:
TELEPHONE: 415-705-8410
TELEPRAX: 415-397-8338
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                 STRANDEDNESS: not relevant
STRANDEDNESS: not relevant
POPOLOGY: not relevant
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-943-120-1
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APPLICANT: HOUSTON, Lou, L.
APPLICANT: SHERIDAN, Philip, J.
APPLICANT: HAWLEY, Stephen
APPLICANT: GLYNN, Jacqueline, M.
APPLICANT: CHAPIN, Steven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19, Application US/09969748C GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 32, Application US/09936759
; GENERAL INFORMATION:
FILING DATE: <Unknown>
                                                                                                                                                                                                                                            LENGTH: 5 amino acids
                            ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: JEFFERSON, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Escherichia coli
US-09-936-759-32
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US-09-936-759-32
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PRICE PRINCIP AND, MARNESH AND METHODS FOR THE TRANSPORT OF BIOLOGICALLY ACTIVE TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TRANSPORT OF BIOLOGICALLY ACTIVE TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR CHANGE AND THE TRANSPORT OF BIOLOGICALLY ACTIVE TITLE OF INVENTION: ACTIVE AND THE TRANSPORT OF BIOLOGICALLY ACTIVE TITLE OF INVENTION: ACTIVE AND THE TRANSPORT OF BIOLOGICALLY ACTIVE THE PRESENCE AND THE TRANSPORT OF BIOLOGICALLY ACTIVE THE PRESENCE AND THE TRANSPORT AND THE TRANSPORT ACTIVE THE TRANSPORT ACTIVE AND THE TRANSPORT OF BIOLOGICALLY ACTIVE THE TRANSPORT OF BIOLOGICAL A
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ORGANISM: Artificial Sequence
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APPLICANT: Kuipers, Maria Eli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100. Matches 1; Conservative
                                                   Conservative
                     Best Local Similarity
Matches 1; Conserv
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US-09-720-278A-2
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  Query Match
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Sequence 1, Application US/09720278A

GENERAL INFORMATION:

APPLICANT: Swart, Pieter Jacob

APPLICANT: Meijer, Dirk Klaas Fokke

APPLICANT: Hageman, Robert John Joseph

TITLE OF INVENTION: Preventing surface infections caused by microorganisms

FILE OF INVENTION: Preventing surface infections caused by microorganisms

FILE REFERENCE: 702-002214

CURRENT APPLICATION NUMBER: US/09/720,278A

CURRENT PILING DATE: 1999-06-28

PRIOR APPLICATION NUMBER: PCT/EP99/04067

PRIOR APPLICATION NUMBER: NL 1010284

PRIOR FILING DATE: 1998-11-06

PRIOR FILING DATE: 1998-10-09

PRIOR FILING DATE: 1998-10-09

PRIOR FILING DATE: 1998-06-26

PRIOR APPLICATION NUMBER: NL 1009505

PRIOR APPLICATION NUMBER: NL 1009505
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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APPLICANT: Lenter, Martin U.
APPLICANT: Lenter, Martin U.
APPLICANT: Beters, Stefan
APPLICANT: Garin-Chea, Pilar
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, Dietmar
APPLICANT: Telan, Leila A.
TITLE OF INVENTION: FAP-Activated Anti-Tumour Compounds
FILE REFERNCE: 1/1087 US
CURRENT APPLICATION NUMBER: US/09/553,800D
CURRENT FILING DATE: 2000-04-21
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 28
SOOTHWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                         Sequence 16, Application US/09553800D
GENERAL INFORMATION:
APPLICANT: Firestone, Raymond A.
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SOFTWARE: Microsoft Word 97 SR-2
SEQ ID NO 1
LENGTH: 5
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ORGANISM: Artificial Sequence
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; OTHER INFORMATION: Precursor
US-09-553-800D-16
                                                                                                   US-09-553-800D-16
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Bet Local Similarity 10.00, 09, Pred. No. 13.54-09, Indels 0, Gaps 0, Match

Matches 1, Conservative 0, Mismatches 0, Indels 0, Gaps 0, De 2 M 2

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BESULT 52

BESULT 53

BESULT 54

BESULT 54

BESULT 55

BESULT
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APPLICANT: Hageman, Robert Johan Joseph
APPLICANT: Wan Den Berg, Jercen Johannes Maria
APPLICANT: Van Den Berg, Jercen Johannes Maria
APPLICANT: Van Den Berg, Jercen Johannes Maria
TITLE OF INVENTION: Pharmaceutical preparations for use in combatting or
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REFERENCE: 702-002214
CURRENT APPLICATION NUMBER: US/09/720,278A
CURRENT PILING DATE: 1999-06-29
FRIOR PILING DATE: 1999-06-29
FRIOR PELING DATE: 1999-11-06
FRIOR PELING DATE: 1998-11-06
FRIOR APPLICATION NUMBER: NL 1010284
FRIOR PELING DATE: 1998-10-09;
FRIOR APPLICATION NUMBER: NL 1009505
FRIOR APPLICATION NUMBER: NL 1009505
FRIOR FILING DATE: 1998-10-06
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APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Waiper, Dirk Klaas Fokke
APPLICANT: Hageman, Robert Johan Joseph
APPLICANT: Wan Den Berg, Jeroen Johannes Maria
APPLICANT: Van Den Berg, Jeroen Johannes Maria
TITLE OF INVENTION: Pharmaceutical preparations for use in combatting or
TITLE OF INVENTION: preventing surface infections caused by microorganisms
FILE REPRENUE: 702-002214
CURRENT APPLICATION NUMBER: US/09/720,278A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Swart, Pieter Jacob
APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Meijer, Dirk Klaas Fokke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 21, Application US/09720278A; GENERAL INFORMATION:
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SOFTWARE: Microsoft Word 97 SR-2
SEQ ID NO 3
                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 1; Conserva
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US-09-720-278A-21
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APPLICANT: Swart, Dieter Jacob
APPLICANT: Wart, Pieter Jacob
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Wan Den Berg, Jeroen Johannes Maria
APPLICANT: Van Den Berg, Jeroen Johannes Maria
APPLICANT: Van Den Berg, Jeroen Johannes Maria
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REFERENCE: 702-002214
CURRENT FILING DATE: 2001-05-24
PRIOR PAPLICATION NUMBER: BP 98203765.7
PRIOR APPLICATION NUMBER: BP 98203765.7
PRIOR PELING DATE: 1998-11-06
PRIOR PLING DATE: 1998-11-06
PRIOR PLING DATE: 1998-10-09
PRIOR PLING DATE: 1998-10-09
PRIOR PLING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 28
SOFTWARRE: Microsoft Word 97 SR-2
SECTION 23
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                 PRIOR FILING DATE: 1999-06-28
PRIOR APPLICATION NUMBER: EP 98203765.7
PRIOR APPLICATION NUMBER: BE 98203765.7
PRIOR APPLICATION NUMBER: NL 1010284
PRIOR FILING DATE: 1998-10-09
PRIOR FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 28
SOSTWARE: Microsoft Word 97 SR-2
SEQ ID NO 21
PRIOR APPLICATION NUMBER: PCT/EP99/04067
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APPLICANT: Swart, Pieter Jacob
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 23, Application US/09720278A; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Matches 1; Conservative
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## APPLICANT: Kuipers, Maria Elizabeth
## APPLICANT: Meijer, Dirk Klaas Fokke
## APPLICANT: Wan Den Berg, Jeroen Johannes Maria
## TITLE OF INVENTION: Preventing Surface infections caused by microorganisms
## TITLE OF INVENTION: Preventing surface infections caused by microorganisms
## TITLE OF INVENTION: Preventing Surface infections caused by microorganisms
## TITLE OF INVENTION: PREVENTION NUMBER: US/09/720,278A
## CURRENT APPLICATION NUMBER: US/09/720,278A
## PRIOR PILING DATE: 1999-06-28
## PRIOR APPLICATION NUMBER: EP 98203765.7
## PRIOR APPLICATION NUMBER: M. 1010284
## PRIOR PILING DATE: 1998-10-09
## PRIOR PILING DATE: 1998-06-26
## NUMBER OF SEQ ID NOS: 28
## SOFTWARE Microsoft Word 97 SR-2
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; OTHER INFORMATION: Description of Artificial Sequence: sequential
; OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-30
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GENERAL INFORMATION:
APPLICANT: WANG, BTYAN S.;
APPLICANT: WANG, BTYAN S.;
ITILE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-USI
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 06/148,422
PRIOR FILING DATE: 1999-08-11
; NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
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ORGANISM: Artificial Sequence
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US-09-636-243B-30
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OTHER INFORMATION: Description of Artificial Sequence: sequential
COTHER INFORMATION: block reoptimization sequence
US-09-636-243B-35
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GENERAL INFORMATION:
APPLICANT: WANG, BTYPE
TITLE OF INVENTION:
FILE REFERENCE:
CURRENT PAPLICATION NUMBER: US/09/636,243B
CURRENT PELING DATE: 1004 / M4-US1
CURRENT PELING DATE: 2000-08-10
PRIOR PELING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN UNINGER: 820
SEQ ID NO 44
SEQ ID NO 44
TYPE: PRT
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US-09-636-243B-56
Sequence 56, Application US/09636243B
GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8225-1004 / M4-US1
CURRENT APPLICATION UNMERR: US/09/636,243B
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR SEQ ID NOS: 83
SOFTWARE: PALENTIN Ver. 2.0
APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-USI1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 35
LENGTH: 5
                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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; OTHER INFORMATION: Description of Artificial Sequence: sequential
; OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-58
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                                                                        OTHER INFORMATION: Description of Artificial Sequence: sequential OTHER INFORMATION: block reoptimization sequence
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47.8%; Score 11; DB 5; Le
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
TITLE OF INVENTION: DIMERIZING PEPTIDES:
FILE REFERENCE: 8125-1004 / M4-US1;
CURRENT APPLICATION VUMBER: US/09/636,243B;
CURRENT FILING DATE: 1999-08-10;
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.
FAPLICANT: WANG, Bryan S.
FILE PEPERENCE: 8325-1004 / M4-USI
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 60148,422
PRIOR APPLICATION NUMBER: 60148,422
FRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VET. 2.0
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ORGANISM: Artificial Sequence
TYPE: PRT
ORGANISM: Artificial Sequence
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                                                 0; Indels
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Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                               SCOURTE SOLUTION OS APPLICATION US/09636243B
GENERAL INFORMATION:
APPLICANT: WANG, BYAN S.
APPLICANT: WANG, BYAN S.
APPLICANT: WANG, BYAN S.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR FULING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO SP
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GENURAL INFORMATION:
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
TITLE OF INVENTION: DIMERIZING FEFTIDES
FILE REPERENCE: 8225-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATCHING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
ISOFTWARE: PATCHIN Ver. 2.0
SEQ ID NO 60
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US-09-636-243B-60
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TYPE: PRT ORGANISM: Artificial Sequence
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US-09-636-243B-76
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OTHER INFORMATION: Description of Artificial Sequence: sequential

OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-62
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                               APPLICANT: WANG, Eryan S.
APPLICANT: WANG, Eryan S.
APPLICANT: PABO, CALI O.
TILE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: WANG, Bryan S.
APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
TITLE REFERENCE: 8325-1004 / M4-US.
CURRENT APPLICATION NUMBER: US/09/636,243B;
CURRENT FILING DATE: 2000-08-10
PRIOR PELING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VET. 2.0
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APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8225-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
US-09-636-243B-61; Sequence 61, Application US/09636243B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-636-243B-62; Sequence 62, Application US/09636243B; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                              SEQ ID NO 61
LENGTH: 5
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OTHER INFORMATION: Description of Artificial Sequence: sequential
OTHER INFORMATION: block reoptimization sequence
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OTHER INFORMATION: Description of Artificial Sequence: sequential

OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-64
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US-09-636-243B-64

Sequence 64, Application US/09636243B

Sequence 64, Application US/09636243B

SEQUENCEAL INFORMATION:
APPLICANT: WANG, Bryan S.
APPLICANT: WASO, CAIL O.
TITLE OF INVENTION: DIMERIZING PEFFIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION UNMBER: US/09/636, 243B

CURRENT FILING DATE: 1999-08-11

NUMBER OF FILING DATE: 1999-08-11

NUMBER OF SEQ ID NOS: 83

SEQUENCE FILING DATE: 200
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
FILE PERERENCE: 8125-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION WUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VEY: 2.0
                   PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 63
LENGTH: 5
CURRENT FILING DATE: 2000-08-10
                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
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## APPLICANT: MAEDA, Hitoaki

## APPLICANT: WAEDA, Hitoaki

## APPLICANT: USHIO, Yoshitaka

## APPLICANT: USHIO, Yoshitaka

## APPLICANT: USHIO, Yoshitaka

## APPLICANT: USHIO, Yoshitaka

## APPLICANT: HITOGUMI, Hitofuui

## APPLICANT: HITOGUMI, Hitofuui

## APPLICANT: NAKATA, Motomized Immunoglobulins Specifically Reactive to Fas Ligand

## TITLE OF INVENTION: Fragments Thereof, and Apoptosis-Induced Site From Fas Ligand

## TITLE OF INVENTION: Fragments Thereof, and Apoptosis-Induced Site From Fas Ligand

## TITLE OF INVENTION: Fragments Thereof, and Apoptosis-Induced Site From Fas Ligand

## TITLE OF INVENTION NUMBER: US/02/24,180C

## CURRENT APPLICATION NUMBER: 271546/1996

## RIOR FILING DATE: 1996-09-20

## PRIOR FILING DATE: 1996-09-20

## NUMBER OF SEQ ID NOS: 183

## SEQ ID NO I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Rucelahti, Erkki
APPLICANT: Resqualini, Renata
APPLICANT: Resqualini, Renata
APPLICANT: Rajotte, Daniel
TITLE OF INVENTION: Methods of Identifying Lung Homing Molecules Using
TITLE OF INVENTION: Membrane Dipeptidase
FILE REFERENCE: P-LA 4377
CURRENT APPLICATION NUMBER: US/09/676,475A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 452
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 278
LENGTH: 5
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                                                                                                                                                                             ; Sequence 1, Application US/09254180C; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 1; Conserva
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ORGANISM: Mouse
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                                                                                                                                RESULT 73
US-09-254-180C-1
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                         OTHER INFORMATION: Description of Artificial Sequence: sequential; OTHER INFORMATION: block reoptimization sequence US-09-636-243B-76
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                                                                                                                                Length 5;
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Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.; APPLICANT: PABO, Carl O.; TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 66/148,422
PRIOR APPLICATION NUMBER: 66/148,422
PRIOR APPLICATION NUMBER: 66/148,422
PRIOR SEQ ID NOS: 83
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: WANG, Bryan S.
APPLICANT: PABO, CAT.1 O.
TILE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-USI
CURRENT APPLICATION WUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-09-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN US: 83
                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 77, Application US/09636243B; GENERAL INFORMATION:
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US-09-636-243B-77
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RESULT 75

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Sequence 1, Application US/09920262A

GENERAL INFORMATION:

APPLICANT: Shallon, Bernie

APPLICANT: Scallon, Bernie

APPLICANT: CANIER CENO. 1

APPLICANT: GILSE-Komar. Jill

PRIOR FILING DATE: 2000-08-07

PRIOR PLING DATE: 2000-08-07

PRIOR PLING DATE: 2000-09-29

PRIOR PLING DATE: 2000-09-29

NUMBER: OF SEQ ID NOS: 15

SECTION 0. 1

LENGTH: S

SECTION 0. 1

LENGTH: S

CORGANISM: How sapiens

US-09-20-262A-1

Query Match

Best Local Similarity 100.0%; Pred. No. 3.56+05;

Matches 1; Conservative 0; Mismatches 0; Indels 0; Gaps

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Search completed: Pebruary 12, 2003, 10:55:52

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US-09-620-412C-9
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US-09-729-597-148
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US-07-657-769B-16
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11, Appl
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            GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 100 summaries
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Gapop 10.0 , Gapext 0.5
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Match Length I
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0; Gaps
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Patent No. 5770377

GENERAL INFORMATION:

APPLICANT: Picksley, Steven M.

APPLICANT: Lane, David P.

TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400

GITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Picksley, Steven M.

APPLICANT: Picksley, Steven M.

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS: ADDRESS: Plehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match

82.6%; Score 19; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FLING DATE: 19-APR-1995
ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UL-1994
ATTORNEY/AGENT INFORMATION:
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Patent No. 5770377
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TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 4:
SEGUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: unknown
US-08-424-957-4
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GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Dicksley, Steven M.
APPLICANT: Dicksley, Steven M.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82.6%; Score 19; DB 1; Length 5; 80.0%; Pred. No. 2e+05; tive 0; Mismatches 1; Indels
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                                                       COMPUTER FRAIABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INPORMATION:
NAME: DEPGE: Malter H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                   NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRI:
ZIP: 94111-418'
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
~~woTTER: IBM PC compatible
...mem: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-APR-1995
UMBER: US 08/277,660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US 08/424,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 4, Application US/09035686
; Patent No. 6153391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/.
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/.
FILING DATE: 20-4UL-1994
ATTORNEY, AGENT INFORMATION:
NAME: Dreger, Walter H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
: California
RY: United States
94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 80.v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: unknown US-08-424-957-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 FXELW 5
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                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-035-686-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 3
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Length 5;
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                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/09099053
Patent No. 6388063
GENERAL INFORMATION:
APPLICANT: Greg Plowman
APPLICANT: David Markby
APPLICANT: Bara Courtneidge
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: SAD RELATED DISORDERS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 69.6%; Score 16; DB 4; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: SUCIAGE SUCIAGE COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FESTERED CON WINDOWS 2.0 CURRENT APPLICATION NUMBER: US/09/099,053 FILING DATE: Herewith CLASSIFICATION DATA: RPLICATION NUMBER: 60/049,914 FILING DATE: June 18, 1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 35" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 235/121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 6
US-07-893-930-37
Sequence 37, Application US/07893930
Patent No. 531/014
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: comessisticated structures of the structure of the 
                                                          1 FXXLW 5
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                                                                                                                                       1 FXELW
                                                                                                                                                                                                                                                                          RESULT 5
US-09-099-053-3
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Sequence 5, Application US/09035686

Patent No. 615391

GENERAL INFORMATION:
APPLICANT: Lane, David P.

TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEB: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Emparcadero Center, Suite 3400
CORRESPONDENCE ADDRESS:
ADDRESSEB: Flehr, Hobbach, Test, Albritton & Herbert
COUNTRY: San Francisco
STREET: Four Emparcadero Center, Suite 3400
COUNTRY: United States
COUNTRY: United States
COMPUTER REABALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: BAPLICATION DATE:
CORPUTER: BAPLICATION DATE:
CORRENT APPLICATION DATE:
CORRENT APPLICATI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              82.6%; Score 19; DB 4; Length 5; 80.0%; Pred. No. 2e+05; tive 0; Mismatches 1; Indels
                               REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTREISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,130
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 78-1989
TELEPR: 910 277299
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
         24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
LENGTH: 5 amino acids
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: amino acid
; STRANDEDNESS:
; TOPOLOGY: unknown
US-09-035-686-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: unknown
US-09-035-686-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 FXXLW 5
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                                                           APPLICANT: Paladino, Joseph
APPLICANT: Bonnet, Jacqueline
APPLICANT: Bonnet, Jacqueline
APPLICANT: Thurieau, Christophe
TITLE OF INVENTION: NEW PEPTIDES
TITLE OF INVENTION: COMPOUNDS DERIVED FROM TACHYKININ
NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS: 46
CORRESPONDENCE ADDRESS: Gordon W. Hueschen
STREET: 715 The "H" Bldg., 310 Bast Michigan
                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: I Eloppy disk
COMPUTER: I Eloppy disk
COMPUTER: I El PR FC COMPALIBLE
COMPUTER: Patent In Release #1.0, Version
SOFTWARE: #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/893,930
FILING DATE: 19920603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 19920603
CLASSIFICATION DATE: 19920603
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: FR 91.06721
FILING DATE: 04.JUN-1991
ATTORNEY, FAGENI INFORMATION:
NAME: Hueschen, Gordon W.
REGISTRATION NUMBER: 16,157
REFERENCE/DOCKET NUMBER: ADIR128/dlk
TELECOMMINICATION INFORMATION:
TELECOMMINICATION INFORMATION:
TELECOMMINICATION OF SEQ. 2030
INFORMATION FOR SEQ ID NO: 37:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
MODISCULE TYPE: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: 1
OTHER INFORMATION: /note= "Benzhydryl
OTHER INFORMATION: carbonyl"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: /note= "NH2"
US-07-893-930-37
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              Kucharczyk, Nathalie
Morris, Angela D.
Fauchere, Jean-Luc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Modified-site
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                                                                                                                                                                                                                                                                                                                            Kalamazoo
                                                                                                                                                                                                                                                                                                                                                                              USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 LW 4
                                                                                                                                                                                                                                                                                                        STREET:
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                                                                                                                                                                                                                                                                                                                                                  STATE:
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Patent No. 5658727
GENERAL INFORMATION:
APPLICANT: Barbas, Carlos
APPLICANT: Barbas, Carlos
APPLICANT: Barbas, Carlos
APPLICANT: Berbas, Carlos
APPLICANT: Lerner: Richard A.
TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING
TITLE OF INVENTION: PHAGEMIDS
NUMBER OF SEQUENCES: 161
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of
ADDRESSEE: Patent Counsel
STREET: 10666 No. 5658727th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
65.2%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                 TITLE OF INVENTION: SYMTHETIC PEPTIDE INHIBITORS OF HIV TITLE OF INVENTION: TRANSMISSION NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS: Pennie & Edmonds STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                            COMPUTER FEADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: EN FC comparible
COMPUTER: EN FC comparible
COMPUTER: Date FOODS MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/073,028
FLING DATE: 07-JUN 1993
CLASSIFICATION: 514
ATTORNEY AGENT INFORMATION:
NAME: COTAZI, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7872-004-999
TEBECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
APPLICANT: Bolognesi, Dani P. APPLICANT: Matthews, Thomas J. APPLICANT: Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: peptide US-08-073-028-44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: unknown
                                                                                                                                                                                                                                             New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 La Jolla
CA
                                                                                                                                                                                                                 CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                          USA
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                                                                                                                                                                                                                                                                                               10036
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US-08-133-011-148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGIH:
                                                                                                                                                                                                                                                                          COUNTRY:
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GENERAL INFORMATION:
APPLICANT: Douvas, Angeline
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
APPLICANT: Takehana, Yoshi
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMINOINPECTIVE CLUSTER VIRUS INFECTIONS
OUNDES OF SEQUENCES: 12.1
CORRESPONDENCE ADDRESS:
ADDRESSER: ROBbins, Berliner & Carson
STREET: 201 No. 5707626th Figueroa Street, Suite 500
CITY: Los Angeles
STREET: California
COUNTRY: U.S.A.
ZIP: 90012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match

65.2%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/704,170
FILING DATE:
                                             APPLICATION NUMBER: DE P 39 04 040.2 FILING DATE: 10-FEB-1989
ATTONNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28957
REFERENCE/DOCKET NUMBER: 18644-96040
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION SEQ 100-962-4810
INFORMATION FOR SEQ 1D NO: 11:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1920-331
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; Patent No. 5707626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Spitals, John P.
REGISTRATION NUMBER: 29,215
REFERENCE/DOCKET NUMBER: 195
TELECOMMUNICATION: TELEPHONE: (213) 977-1001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 103:
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LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                   LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                            ) MOLECULE TYPE: peptide
; HYPOTHETICAL: YES
US-08-584-579-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: peptide US-08-704-170-103
                             PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           linear
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3Y: linear
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; Patent No. 5674979
; GENERAL INFORMATION: APPLICANT: Schramm, Wolfgang
APPLICANT: Schramm, Hans J.
; TITLE OF INVENTION: Agent For Inhibiting Symmetrical
TITLE OF INVENTION: Proteins, In Particular Enzymes
NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
ADDRESSE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 1; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
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ZIP: 20005

MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,579
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/332,447
FILING DATE: 31-OCT-1994
APPLICATION NUMBER: US 08/112,215
FILING DATE: 26-AUG-1993
APPLICATION NUMBER: US 07/976,003
FILING DATE: 13-NOV-1992
APPLICATION NUMBER: US 07/585,141
FILING DATE: 07-DEC-1990
APPLICATION NUMBER: US 07/585,141
FILING DATE: 07-DEC-1990
APPLICATION NUMBER: US 07/585,141
FILING DATE: 07-DEC-1990
APPLICATION NUMBER: US 07/585,141
FILING DATE: 09-FEB-1990
FILING DATE: 08-JUN-1994
CLASSIFICATION: 435
PRIOR APPLICATION BATA:
APPLICATION NUMBER: US 07/683,602
FILING DATE: 10-APR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/826,623
FILING DATE: 27-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US 92/03091
FILING DATE: 10-APR-1992
ATTORNEY/AGENT INFORMATION:
NAMME: FILLING DATE: 34,163
REFERENCE/DOCKET NUMBER: SCRF 238.2
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: 619-554-6312
INFORMATION FOR SEQ ID NO: 148:
SEQUENCE CHARACTERIFICS:
LENGTH: 5 amino acids
TYPE: amino acid
TYPE: amino acid
TYPE: mine acid
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Matches 2; Conservative
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COUNTRY: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-133-011-148
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APPLICANT: HILGER, CHRISTOPH-STEPH
APPLICANT: DINKELBORG, LUDGER
APPLICANT: KRAMP, WOLGANG
APPLICANT: SCHIER, HANS-WARTIN
TITLE OF INVENTION: CHELATORS OF TYPE XNISIOI FOR RADIOACTIVE ISOTOPES, 7H
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 25,329
REPERENCE/DOCKET VUMBER: WBLT 2006 (WO) PA
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-362-3800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/08854187A
Patent No. 5562642
GENERAL INFORMATION:
APPLICANT: TISTE OF INVENTION: Method for Sequencing of TITLE OF INVENTION: Protein or Peptide
                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: FLIESLER, DUBB, MEYER & LOVEJOY
STREET: Four Embarcadero Center, Suite 400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. wc.
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FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Dubb, Hubert E
REGISTRATION NUMBER: 25,329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT/DE94/00372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
SOFTWART APPLICATION DATA:
APPLICATION NUMBER: US/08/530,326
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Adams & Wilks
STREET: 50 Broadway, 31st Floor
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: Diskette COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTHARE: FASTER: THE PASTER OF TH
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/D
FILING DATE: 29-MAR-1994
APPLICATION NUMBER: P 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 415-362-380
TELEFAX: 415-362-2928
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 2; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amino acid
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         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FRAGMENT TYPE: i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE
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US-08-854-187A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Ne COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
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CORRESPONDENCE ADDRESS: ADDRESS: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, 8th Floor CITY: San Francisco STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 2; Length 5; ilarity 100.0%; Pred. No. 2e+05; Conservative 0; Mismatches 0; Indels
      Query Match 65.2%; Score 15; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30
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APPLICATION NUMBER: US/08/747,137
FILING BATE: 12-NOV-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION DATA:
FILING DATE: 14-NAR-1994
FRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/069,831
FILING DATE: 01-UN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/959,560
FILING DATE: 13-OCT-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US 07/641,720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 12
US-08-530_326-14
; Sequence 14, Application US/08530326
; Patent No. 5961954
                                                                                                                                                                                                                                                                                                                                                Sequence 44, Application US/08747137 Patent No. 5945033 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Apple, Randolph T.
REGISTRATION UNDRER: 36,429
REFRENCE/DOCKET UNDRER: 01615
TELECOMMUNICATION INFORMATION:
TELEHONE: 415-576-0200
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 0° FILLING DATE: 15-JAN-1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 amino acids
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Matches 2; Conserva
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RESULT 15
US-08-862-625A-3
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TITLE OF INVENTION: MONOSPECIFIC ANTIBODY REACTIVE WITH
TITLE OF INVENTION: MATRIX METALLOPROTEINASE CLEAVAGE PRODUCTS OF FIBRIN(OGEN)
NUMBER OF SEQUENCES:
ADDRESSEE: Hoffmann & Baron, LLP
STREET: 350 Jericho Turnpike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 65.2%; Score 15; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
COMPUTER READBLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: NEC PowerMate VP75
COPTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/854,187A
FILING DATE: 09-MAY-1997
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: JP 08-115087
FILING DATE: 09-MAY-1996
APPLICATION NUMBER: JP 08-154580
FILING DATE: 1996
APPLICATION NUMBER: JP 08-154580
FILING DATE: 1996
APPLICATION NUMBER: JS,386
RECISTRATION NUMBER: SS,386
RECISTRATION NUMBER: SS,386
REFERENCE/DOCKET NUMBER: SS,004-3185
TELEGOMMUNICATION INFORMATION:
TELEFAX: (212) 809-3700
TELEFAX: (212) 809-3700
TELEFAX: CHARACTERISTICS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/900,895
FILING DATE: 7/25/97
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: O'DEA, Sean W.
REGISTRATION NUMBER: 37690
REGISTRATION NUMBER: 454-16
TELECHONE: (516) 822-3550
TELECHONE: (516) 822-3550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/08900895
Patent No. 6043087
GENERAL INFORMATION:
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STRANDEDNESS: not applicable
TOPOLOGY: linear
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STATE: New York
COUNTRY: USA
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65.2%; Score 15; DB 3; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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; Sequence 3. 604603
; GENERAL INFORMATION:
    APPLICANT: Tsugiata, Akira
    TITLE OF INVENTION: A method for Amino Acid
    TITLE OF INVENTION: Sequencing of Protein or
    TITLE OF INVENTION: Peptide from Carboxy Terminus
    TITLE OF INVENTION: Peptide from Carboxy Terminus
    TITLE OF INVENTION: Thereof
    NUMBER OF SEQUENCES: 3
    CORRESPONDENCE ADDRESS:
    ADDRESSE: Adams & Wilks
    STREET: 50 Broadway, 31st Floor
    CHTY: New YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette, 3.50 inch, MEDIUM TYPE: 1.44 Mb storage COMPUTE: IBM PS/2 OPERATING SYSTEM: DOS 3.3 SOFTWARE: Wordperfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/862,625A FILING DATE: 24-MAY-1997 CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 24-MAY-1997
CLASSIFICATION MAY-1997
CLASSIFICATION 1436
PRIOR APPLICATION DATE: 34-MAY-1997
FILING DATE: 24-MAY-1996
PRIOR APPLICATION DATE: 49-MAY-1996
PRIOR APPLICATION DATA: APPLICATION NUMBER: JP 09-035312
FILING DATE: 19-FEBRUARY-1997
PRIOR APPLICATION NUMBER: JP 09-036610
FILING DATE: 20-FEBRUARY-1997
PRIOR APPLICATION DATA: APPLICATION NUMBER: JP 09-091215
FILING DATE: 09-APPLICATION NUMBER: JP 09-091215
FILING DATE: 09-APPLICATION NUMBER: JP 09-091215
FILING DATE: 09-APPLICATION:
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REGISTRATION NUMBER: 25,386
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COUNTRY: United States of America
ZIP: 10004
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STRANDEDNESS: not applicable
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 809-3704
INFORMATION FOR SEQ ID NO: 3:
                      SEQUENCE CHARACTERISTICS:

LENGTH: 5 amino acids

TYPE: amino acid
TOPOLOGY: linear

MOLECULE TYPE: peptide
US-08-900-895-1
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INFORMATION FOR SEQ ID NO:
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MOLECULE TYPE: peptide
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OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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                                                                                                                                                                                       0; Indels
                                                                                                          Query Match 65.2%; Score 15; DB 3; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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; Patent No. 6127339
; GENERAL INFORMATION:
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; Sequence 51, Application US/08981122B
; Patent No. 6127339
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ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
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                ; OTHER INFORM
US-08-981-122-38
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SEQ ID NO 51
LENGTH: 5
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LENGTH: 5
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OTHER INFORMATION: Sequence of a peptide synthesized in Examples 11 and 12 from L-fd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 6127339 OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid
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OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
OTHER INFORMATION: synthesizing system (RaMPS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: Sequence of a peptide synthesized in Example 2 from L-form
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APPLICANT: Aritomi, Masaharu
TITLE OP INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT PILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: UP 7-176904
PRIOR FILING DATE: 1995-06-21
PRIOR FILING DATE: 1995-06-21
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APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
                                                                                     0; Gaps
Query Match 65.2%; Score 15; DB 3; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: UP 7-176904
PRIOR FILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR APPLICATION NUMBER: PCT/JP96/01734
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PAtentIn Ver: 2.0
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SOFTWARE: Patentin Ver. 2.0
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NAME/KEY: AMIDATION
LOCATION: 5
COCATION: 5
OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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APPLICANT: Hatenaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
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FRAGMENT TYPE: 1
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        US-08-530-342A-14
                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-208-277-9
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                     LOCATION: 5
CTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for Patent No. 6127339
CTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid CTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:

APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Wild, Carl T.
ITIE OF INVENTION: SYNTHETIC PEPTIDE INHIBITORS OF HIV TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPOTER: PACENTION DATA:
APPLICATION NUMBER: US 08/073,028
FILING DATE: 07-UN-1993
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: USA
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US-08-554-616-44
; Sequence 44, Application US/08554616
; Patent No. 6133418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : (212) 790-9090
(212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Coruzzi, Laura A. 742
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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NAME/KEY: AMIDATION LOCATION: 5
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RESULT 21

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Sequence 9, Application US/09208277
Petent No. 6166177
GENERAL INFORMATION:
APPLICANT: Probst, Peter
APPLICANT: Bhatia, Ajay
APPLICANT: Shatia, Ajay
APPLICANT: Skeiky, Yasir A, W.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE TREATMENT AND DIAGNOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                            APPLICANT: DINKELBORG, LUDGER
APPLICANT: DINKELBORG, LUDGER
APPLICANT: KARAMP, WOLGANG
APPLICANT: KRAMP, WOLGANG
APPLICANT: SCHIER, HANS-MARTIN
TITLE OF INVENTION: TYPE S3N2 CHELATORS FOR RADIO
TITLE OF INVENTION: TYPE S3N2 CHELATORS FOR RADIO
TITLE OF INVENTION: THEIR DIAGNOSTIC AND THERAPEUTICAL USE
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                      ADDRESSEE: WOOD, PHILLIPS, VAN SANTEN, CLARK & ADDRESSEE: MORTIMER
                                                                                                                                                                                                                                                                                                                                                                          STREET: 500 West Madison Street, Suite 3800 CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 60661-2511
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: COMPAQ ProLinea 4/25s
OPERATING SYSTEM: DOS
SOFTWARE: WordPerfect for Windows 6.0a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/530,342A
FILING DATE: 08-MAR-1996
CLASSIFICATION: 424
FRICK APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00370
FILING DATE: 39-MAR-1994
APPLICATION NUMBER: P 43 11 022.3
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1214.00008
; Sequence 14, Application US/08530342A; Patent No. 6143275; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Mortimer, John S. REGISTRATION NUMBER: 30,407 REFERENCE/DOCKET NUMBER: 121 TELECOMMUNICATION: 312/876-1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Matches 2; Conservative
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TYPE: amino acid
STRANDEDNESS: single
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MOLECULE TYPE: peptide
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APPLICANT: Barbas, Carlos
APPLICANT: Lerner, Richard
APPLICANT: Lerner, Richard
APPLICANT: HETERODIMERIC RECEPTOR LIBRARIES USING
TITLE OF INVENTION: PHAGEMIDS
                                                                                                                                                                                                                                                                                   Query Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 161
CORRESPONDENCE ADDRESS:
ADDRESSE: The Scripps Research Institute, Office of ADDRESSE: Patent Counsel
STREET: 10666 No. 6235469th Torrey Pines Road, TPC-8
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
TITLE OF INVENTION: OF CHLAMYDIAL INFECTION FILE REFERENCE: 210121.469
CURRENT APPLICATION NUMBER: US/09/208,277
CURRENT FILING DATE: 1998-12-08
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/826,623
FILING DATE: 27-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US 92/03091
FILING DATE: 10-APR-1992
ATTORNEY/AGENT INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/907,739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 148, Application US/08907739; Patent No. 6235469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/133,011
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REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: SCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-554-2937
TELEPRAK: 619-554-6312
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                                                                                                                                                                                           TYPE: PRT ORGANISM: Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CA
USA
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patent No. 6348568

general information:
patent No. 6348568

general information:
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Query Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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65.2%; Score 15; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Anwer, Mohmed APPLICANT: Lambert, Dennis TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED TITLE OF INVENTION: PHARMACOKINETIC PROPERTIES FILE REFERENCE: 7872-043
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: 7872-043
CURRENT APPLICATION NUMBER: US/09/082,279B
CURRENT FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 791, Application US/09082279B
Patent No. 6289782
; GENERAL INFORMATION:
PAPLICANT: Barney, Shawn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTHER INFORMATION: Core polypeptide US-09-082-279B-791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Core polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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Merutka, Gene
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US-09-315-304B-791
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US-09-082-279B-791
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SEQ ID NO 791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
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APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURESH M. DESAI
APPLICANT: THOMAS P. LEARY
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: SHEER
APPLICANT: SHEER IL. BUJNK
APPLICANT: SHEER IL. BUJNK
APPLICANT: SIA K. MUSHAHWAR
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
             GENERAL INFORMATION:
APPLICANT: Steven P. Fling
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121.469C7
CURRENT APPLICATION NUMBER: US/09/620,412C
CURRENT FILING DATE: 2000-07-20
NUMBER OF SEQ ID NOS: 363
SOFTWARE: FASEUSEQ for Windows Version 3.0/4.0
                                                                                                                                                                                                                                                                                             Query Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/469,260A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
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TAMI J. PILOT-MATIAS
GEORGE J. DAWSON
GEORGE G. SCHLAUDER
SURESH M. DESAI
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. C. ERKER
SHERI L. BUIJK
ISA K. MUSHAHWAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-469-260A-228
Sequence 228, Application US/08469260A
Parent No. 6451578
GENERAL INFORMATION:
                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-620-412C-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
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ZIP: 60064-3500
   Patent No. 6448234
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                        4 LW 5
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                                                                                                                                                                                   SEQ ID NO 9
LENGTH: 5
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APPLICANT: Skeiky, Yaair
APPLICANT: Skeiky, Yaair
APPLICANT: Fling, Stewey
APPLICANT: Fling, Stewey
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
FILE REPRESENCE: 201021.469C5
CURRENT APPLICATION NUMBER: US/09/556,877
CURRENT PILING DATE: 2000-04-19
NUMBER OF SEQ ID NOS: 305
SOFTWARE: FastSEQ for Windows Version 3.0/4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       65.2%; Score 15; DB 4; Length 5; 100.0%; Pred. No. 2e+05; Aztive 0; Mismatches 0; Indels
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 0; Indels
 0; Mismatches
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                                                                                                                                                            Sequence 9, Application US/09556877
Patent No. 6432916
GENERAL INFORMATION:
APPLICANT: Probst, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-556-877-9
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Conservative
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Best Local Similarity
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Best Local Similarity
Matches 2; Conserv
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US-09-620-412C-9
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US-09-556-877-9
                                  4 LW 5
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Matches
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Lerner, Richard A.
TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING
PHAGEMIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent Counsel
STREET: 10666 No. 6468738th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: The Scripps Research Institute, Office of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: 08/09/729,597
FILING DATE: 04-Dec-2000
CLASSIFICATION NUMBER: 08/133,011
APPLICATION DATA:
APPLICATION NUMBER: 08/133,011
FILING DATE: 10-409-29
APPLICATION NUMBER: US 07/826,623
FILING DATE: 10-APR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: SCRF 238.2
TELEBPHONE: 619-554-2937
                                                                                  Query Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 4; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                               Sequence 148, Application US/09729597
Patent No. 6468738
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 148:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kang, Angray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
TOPOLOGY: linear
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; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-469-260A-228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: La Jolla
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Best Local Similarity
Matches 2; Conserva
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                                                                                                                                                                                                                                                                            RESULT 30
US-09-729-597-148
                                                                                                                                                                     4 LW 5
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Gaps
                                                                                    TITLE OF INVENTION: Altered Linolenic and Linoleic Acid Content
TITLE OF INVENTION: in Plants
NUMBER OF SEQUENCES: 72
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
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APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPEUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 65.2%; Score 15; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA:
APPLICATION NORSE: PCT/US94/01321
FILING DATE: 04-FEB-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT-US94-02631-103; Sequence 103, Application PC/TUS9402631; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                          FILING DATE: 04-FEB-1994
CLASSIFICATION:
PRIOR APPLICATION NUMBER: US 08/156551
FILING DATE: 22-NOV-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US 08/014431
PCT-US94-01321-47; Sequence 47, Application PC/TUS9401321; GENERAL INFORMATION:
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                             IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 05-FEB-1993
INFORMATION FOR SEQ ID NO: 4'
                                                                                                                                                                                       Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5 amino acids
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MOLECULE TYPE: peptide
PCT-US94-01321-47
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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                                                                                                                                                                                                                                                      Score 14; DB 1; Length 5; Pred. No. 2e+05; 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPEDUDENCE ADDRESS:
ADDRESSE: MORRISON & FOERSTER
STREET: SOOO Pennsylvania Ave., NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
SOFTWARE PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/475,263
FILING DATE: O'-JUN-1995
CLASSIFICATION: 424
ATTONENY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,959
TELEPROMUNICATION INFORMATION:
TELEPROMUNICATION INFORMATION:
TELEPRAKE (202) 887-0763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 83, Application US/08475263; Patent No. 5759994; GENERAL INFORMATION:
                         TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: AMINO ACID
          TELECOMMUNICATION INFORMATION:
                                                                                            83:
                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 40.0%;
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 2006-1812
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 40.0
Matches 2; Conservative
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TELEX: 90-4030
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                     ESS: single
linear
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                                                                                                                                                   TYPE: AMINO A STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                        1 FXXLW 5
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US-07-789-184-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-475-263-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-475-263-83
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 36
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| Patent No. 5688768
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: SCAEBCROUGH, ROBERT M.
| TILLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND TITLE OF INVENTION: RELATED PHARMACEUTICALS
| NUMBER OF SEQUENCES: 223
| CORRESPONDENCE ADDRESS: ADDRESSEE: MORRISON & FOERSTER STREET: 755 Page Mill Road CITY: Palo Alto
| STATE: California COUNTRY: USA
                                                                                                                                                                                                                                                    65.2%; Score 15; DB 5; Length 5; 100.0%; Pred. No. 2e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              65.2%; Score 15; DB 6; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/789,184
FILING DATE: 19911107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 5171838

Patent No. 5171838

TITLE OF INVENTION. LEUJA BINDING PEPTIDES

NUMBER OF SEQUENCES: 24

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/526,921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: WIRASHIES, KATE H.
REFERENCE/DOCKET NUMBER: 29.59
REFERENCE/DOCKET NUMBER: 22000-20502.20
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEPAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 103:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: 1 TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                     ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US94-02631-103
                                                                                                                                                                                                                                                                         Best_Local Similarity
Matches 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 2; Conserv
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US-07-789-184-83
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5171838-3
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                                                                                                                                                         APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RELOADED PHARMACEUTICALS
TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
CORRESPONDENCES: 223
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURREALING SISIEM:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,886
FILING DATE: US/08/485,886
FILING DATE: US/08/485,886
FILING DATE: US/08/485,886
FILING DATE: US/07/789,184
FILING DATE: US/07/789,184
FILING DATE: US/07/189,184
FILING DATE: US/07/189
FILING DATE: US/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                Sequence 83, Application US/08485886
Patent No. 5798248
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 83, Application US/08477362; Patent No. 5849507; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: MORRISON & FOERSTER STREET: 755 Page Mill Road CITY: Palo Alto STATE: California COUNTRY: USA ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-485-886-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 FXXLW 5
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US-08-477-362-83
US-08-485-886-83
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 83, Application US/08477134
Patent No. 585648
GENERAL INFORMATION: SHAUN R.
APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROWBIN RECEPTOR AND TITLE OF INVENTION: RELATED PHARMACEUTICALS
                   SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PLICATION NUMBER: US/08/477,362
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,134
FILING DATE: 07-JUN-1995
CLASSIFCATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
                                                                                                                                                                                                                                                                            22000-20502.20
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                                                                                       FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOBRSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                        NAME: MURSHICE, KATE H.
REGISTRATION NUMBER: 29,959
REFRENCE/DOCKET NUMBER: 22000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 29,959
REFRENCE/DOCKET NUMBER: 2200C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match
Best Local Similarity 40.0%
Lag 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: linear
US-08-477-362-83
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LENGTH: 5 amino acids
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                                                                                                                                                             STREET: 755 Page M
CITY: Palo Alto
STATE: California
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        GENERAL INFORMATION:
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US-08-485-695-83
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                                                                                                                                                                                                                                                                                                                          Sequence 83, Application US/08473489A

Patent No. 6024936

GENERAL INFORMATION:
APPLICANT: COUGHLIN, SHAUN R.
TITLE OF INVENTION: RELATED PHARMACEUTICALS
ITLE OF INVENTION: RELATED PHARMACEUTICALS
ORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
COUNTY: USA
                                                                                                                  60.9%; Score 14; DB 2; Length 5;
40.0%; Pred. No. 2e+05;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 60.9%; Score 14; DB 3; Length 5; Best Local Similarity 40.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: DATE COMPATIBLE
COMPUTER: DATE COMPATIBLE
COMPUTER: DATE CATION DATA:
APPLICATION NUMBER: US/08/473,489A
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/789,184
FILING DATE: 1991-11-07
ATTORNEY/AGENT INFORMATION:
NAMB: MURASHIGE: KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 29,959
REFERENCE/OCKET NUMBER: 22,000-20502.20
TELECOMMUNICATION:
MEDIUM TATION NUMBER: 20,959
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-485-695-83
; Sequence 83, Application US/08485695
; Patent No. 6124101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (415) 813-5600
TELEFAX: (415) 494-0792
                                                                                                              Query Match
Best Local Similarity 40.0°
Matches 2; Conservative
; LENGTH: 5 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-477-134-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                                             1 FXXLW 5
                                                                                                                                                                                                                                  1 FEPFW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-473-489A-83
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APPLICANT: COUGHING, SHANK,

APPLICANT: COUGHING, SHANK,

TITLE OF INVERTION: RECORDER M.

ANDRESSER. COMPRESS. 23

ANDRESSER. COMPRESS. 23

ANDRESSER. CALIFORNIA

CONTRY: To Palo Alto
STATE: To Spee Mill Road

CONTRY: To Palo Alto
STATE: To Spee Mill Road

CONTRY: CALIFORNIA

CONTRIBUTION

CONTRY: CALIFORNIA

CALIFORNIA

CONTRIBUTION

C
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/07960636B; Patent No. 5369015
; INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (202) 659-2930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (202) 887-0357
TELEFAX: (202) 887-0357
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                          1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                 TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                      Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20006
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-07-960-636B-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                       US-07-858-842-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 LW 5
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APPLICANT: Yoshikawa, Kelichi
APPLICANT: Yoshowa, Kelichi
APPLICANT: Hassegawa, Masayasu
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, Fujita, Sergiban For Roducing AN ANGIOTENSIN
NUMBER OF INVENTION: METHOD FOR PRODUCING AN ANGIOTENSIN
SUMBER OF SEGURACES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 60.9%; Score 14; DB 4; Length 5; 40.0%; Pred. No. 2e+05; 1ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/858,842
FILING DATE: 19920327
CLASSITCATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/858,842
FILING DATE: 29-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UP 142283
FILING DATE: 29-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UP 298060
FILING DATE: 17-OCT-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                 22000-20502.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: ARMSTRONG & KUBOVCIK
STREET: 1725 K Street N.W. Suite 1000
                FILING DATE: 17 FEB-1933
CLASSTFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
FILING DATE: 07 NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: MURABHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REGISTRATION NUMBER: 29,959
RECISTRATION NUMBER: 29,959
TELEPHONE: (415) 494-0792
TELEPHONE: (415) 494-0792
TELEFAX: (415) 494-0792
TELEFAX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Wasington
STATE: District of Columbia
COUNTRY: United States of America
  APPLICATION NUMBER: US/08/018,760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      920247
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Patent No. 5314807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 34, 409
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REGISTRATION NUMBER: 34, 409
REFERENCE/DOCKET NUMBER: 920
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 659-2930
TELEFAX: (202) 887-0357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 40.09
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                   LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear
US-08-018-760-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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TELEFAX: (2
TELEX: 4401
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APPLICANT: Yoshikawa, Masaaki
APPLICANT: Yoshikawa, Masaaki
APPLICANT: Yoshikawa, Masaaki
APPLICANT: Yasumoco, Ryouichi
APPLICANT: Yasumoco, Ryouichi
APPLICANT: Yasumoco, Ryouichi
TITLE OF INVENTION: Method for Producing an Angiotensin
TITLE OF INVENTION: Converting Enzyme Inhibitor-Containing Composition
VOMBER OF SEQUENCES:
ADDRESSEE: Armstrong, Westerman, Hattori, McLeland &
ADDRESSEE: Armstrong, Westerman, Hattori, McLeland &
STREET: 1725 K Street, N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/960,636B
FILING DATE: 14-OCT-1992
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: JP 298061/1991
FILING DATE: 17-OCT-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Tockman, Albert
REGISTRATION NUMBER: 19722
REFERENCE/DOCKET NUMBER: P1161-2679-A920798
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Mammalian meat, fish, crustaceans
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US-07-802-667-19
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LOCATION: 5
OTHER INFORMATION: /label= Val-resin
OTHER INFORMATION: /note= "The carboxy terminal amino acid, Val, is
OTHER INFORMATION: attached to a resin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; ative 1; Mismatches 0; Indels
                                                                                             Sequence 8, Application US/08384680
Fatent No. 5464935
GENERAL INFORMATION:
APPLICANT: Hearner, George A.
APPLICANT: Hearner, Douglas
APPLICANT: Miljenko
ITILE OP INVENTION: Peptide Inhibitors of Selectin
ITILE OP INVENTION: Binding
WUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: FIDEPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/384,680
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CCOR-0024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION: 514
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 891,986
FILING DATE: 05-MAY-1992
ATTORNEY, AGENT INFORMATION:
NAME: Elderkin, Dianne B.
REGISTRATION NUMBER: 28,598
REFERENCE/DOCKET NUMBER: CCR-00:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-133-020-4; Sequence 4, Application US/08133020; Patent No. 5476840; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 215-568-3439
TELEX: 710-670-1334
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 215-500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                            STATE: Pennsylvania
COUNTRY: USA
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Best Local Similarity
                                                                                  US-08-384-680-8
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1 IW 2
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APPLICANT: BIHOVSKY, RON H.
APPLICANT: ERHARDT, PAUL W.
APPLICANT: LAMPE, JOHN W.
APPLICANT: MOHAN, RAJU
APPLICANT: SIAW, KENNETH J.
TITLE OF INVENTION: INHIBITORS OF THE CONVERSION OF BIG
TITLE OF INVENTION: ENDOTHELIN TO ENDOTHELIN
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERLEX LABORATORIES, INC.
STREET: 3300 FAIRFIELD ROAD
                            DERIVATIVES, PREPARATION, METHOD AND APPLICATION AS MEDICAMENTS OF SAID NEW DERIVATIVES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        56.5%; Score 13; DB 1; Length 5; ilarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
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ZIP: 07470-5472
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/802,667
                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: ISM PC COMPATIELE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORD PERFECT # 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,020
FILING DATE: 03-DEC-1993
CLASSIFICATION: 530
TITLE OF INVENTION: NEW HEXAPEPTIDIC TITLE OF INVENTION: DERIVATIVES, PREI TITLE OF INVENTION: APPLICATION AS MITITLE OF INVENTION: DERIVATIVES NUMBER OF SEQUENCES: 7 CORRESPONDENCE ADDRESS: ADDRESSE: BIERMAN AND MUSERLIAN STREET: 600 THIRD AVE.
       NEW HEXAPEPTIDIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 19, Application US/07802667; Patent No. 5504070; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-661-8000
TELEFAX: 212-661-8002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19,683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: CHARLES A. MUSERLIAN
REGISTRATION NUMBER: 19,683
REFERENCE/DOCKET NUMBER: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 212-661-8002
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
DESCRIPTION: PEPTIDE
US-08-133-020-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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LENGIH: 5 amino acids
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APPLICANT: ERHARDY, PAUL W.
APPLICANT: LAMPE, JOHN W.
APPLICANT: CAMPE, JOHN W.
APPLICANT: SHAW, KAUN
APPLICANT: SHAW, KENNETH J.
TITLE OF INVENTION: INHIBITIONS OF THE CONVERSION OF BIG
TITLE OF INVENTION: INHIBITIONS OF THE CONVERSION OF BIG
TITLE OF INVENTION: SUDOTHELIN
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERLEX LABORATORIES, INC.
STREET: 3300 FAIRFIELD ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 56.5%; Score 13; DB 1; Length 5; Best Local Similarity 50.0%; Pred. No. 2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: 5
; COTHER INFORMATION: /note= "C-TERMINAL AMIDE ESTER"
US-07-802-667-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT PAPLICATION DATA:
APPLICATION NUMBER: US/07/802,667
FILING DATE: 19911205
CLASSIFICATION: 51
ATTORNEY/AGENT INFORMATION:
NAME: BELLAMY, ELIZABETH A.
REFERENCE/DOCKET NUMBER: BERLEX 51
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL'
                                                                                                                                  BERLEX 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20, Application US/07802667 Patent No. 5504070 GENERAL INFORMATION:
                      CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: BELLAMY, ELIZABETH A.
REGISTRATION NUMBER: 27,854
REFERENCE/DOCKET NUMBER: BERLE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-305-4405
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 07470-5472
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Modified-site LOCATION: 5
                                                                                                                                                                                                                                                                                         LENGTH: 5 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   201-305-4405
FILING DATE: 19911205
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -07-802-667-20
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TITLE OF INVENTION: ENDOTHELIN TO ENDOTHELIN
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERLEX LABORATORIES, INC.
STREET: 300 FAIRFIELD ROAD
CITY: WAYNE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
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ZIP: 07470-5472
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BM PC compatible
OMPOTER: IBM PC compatible
OMPRANIE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/802,667
FILING DATE: 19911205
ATCASSIFICATION: ELIZABETH A.
ATCASSIFICATION NUMBER: 27,854
ATCASSIFICATION NUMBER: 27,854
ATCASSIFICATION NUMBER: 27,854
FREFRENCE/DOCKET NUMBER: BERLEX 51
TELEGHONE: 201-305-5472
TELEGHONE: 201-305-5472
TELEGHAX: 201-305-5472
TELEGHAX: 201-305-5472
TELEGHAX: 201-305-305-5472
TELEGHAX: 201-305-305-5472
TELEGHAX: SEQUENCE CHARACTERISTICS:
FROUGHUE CHARACTERISTICS:
FROUGHUE SEQUENCE:
FROUGHUE CHARACTERISTICS:
FROUGHUE CHARACTERISTICS
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OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL"
                                                                                                                                                                                                                                                              LOCATION: 1
OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21, Application US/07802667; Patent No. 5504070
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BIHOVSKY, RON H.
APPLICANT: BRHARDT, PAUL W.
APPLICANT: LAMPE, JOHN W.
APPLICANT: MOHAN, RAJU
APPLICANT: SHAW, KENNETH J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Modified-site
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                                                                                                                                                                                                                 NAME/KEY: Modified-site
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Matches 1; Conservative
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                   MOLECULE TYPE: peptide FEATURE:
TYPE: AMINO ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide
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                                                                                         linear
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                                                                                   Query Match 56.5%; Score 13; DB 1; Length 5; Best Local Similarity 50.0%; Pred. No. 2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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; LOCATION: 5
; OTHER INFORMATION: /note= "C-TERMINAL AMIDE ESTER"
US-07-802-667-21
                                                                                                                                                                                                                                                                      RESULT 49
US-08-057-167-12
i Sequence 12, Application US/08057167
j Patent No. 5541095
i GENERAL INFORMATION:
APPLICANT: Hischberg, Carlos B.
APPLICANT: Hashimoto, Yasuhiro
APPLICANT: Swiedler, Stuart J.
APPLICANT: Swiedler, Stuart J.
APPLICANT: Swiedler, Stuart J.
APPLICANT: Ishihara, Masayuki
TITLE OF INVENTION: GLYCOSAMINOGLYCAN SPECIFIC
TITLE OF INVENTION: SULFOTRANSFERASES
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: Massachusetts
CONNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER: U.S.A.
COMPUTER: BADABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: TBM PS/2 Model 502 or 555X
OPBERATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/057,167
FILING DATE: 19930430
CLASSIFICIATION 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/899,432
FILING DATE: 16 June 1992
ATTORNEY/AGENT INFORMATION:
NAME: CLARK, PAUL T.
REGISTRALION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 04020/015002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B: Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 200154
INFORMATION FOR SEQ 1D NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 5
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US-08-057-167-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boston
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STREET: 22
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US-08-269-257-9
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              Parent No. 5550110
GENERAL INFORMATION:
APPLICANT: Cody, wayne L.
APPLICANT: Topliss, John G.
TITLE OF INVENTION: Endothelin Antagonists II
NUMBER OF SEQUENCES:
ADDRESSEE: Warner-Lambett Company
STREET: 2800 Plymouth Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5, Application US/08373911
Patent No. 5663296
GENERAL INFORMATION:
APPLICANT: Doherty, Annette M.
APPLICANT: Kornberg, Brian E.
APPLICANT: Nikam, Sham S.
ITILE OF INVENTION: Hydroxamate Inhibitors of
ITILE OF INVENTION: Endothecin Converting Enzyme
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Ver. #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/269,257
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CLASSIFICATION: 514

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PETLING DATE: 31-MAR-1993

ATTORNEY AGENT INFORMATION:

NAME: Tinney, Francis J.

REGISTRATION NUMBER: 9D-4334-P2

REFERENCE/DOCKET NUMBER: 9D-4334-P2

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEPHONE: 313 996-1295

TELEFAX: 313 996-1253

INFORMATION FOR SEQ ID NO: 9:

SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                               ZIP: 48105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
Sequence 9, Application US/08269257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 48105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 2800 Plymouth Rd CITY: Ann Arbor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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US-08-373-911-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Doherty, Annette M.
APPLICANT: Kornberg, Brian E.
APPLICANT: Nikam, Sham S.
TITLE OF INVENTION: Hydroxamate Inhibitors of
TITLE OF INVENTION: Endothecin Converting Enzyme
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRET APPLICATION DATA:
APPLICATION NUMBER: US/08/373,911
                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/070,161
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Warner-Lambert Company
STREET: 2800 Plymouth Rd.
CITY: Ann Arbor
                                                                                  FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/070,161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Tinney, Francis J.
REGISTRATION NUMBER: 33069
REFERENCE/DOCKET NUMBER: PD-4435
TELECOMMUNICATION INFORMATION:
TELEPHONE: 313 996-7295
                                                                                                                                                                                                                                                                  PD-4435
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                   FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Tinney, Francis J.
REGISTRATION NUMBER: 33069
REFERENCE/DOCKET NUMBER: DD-44:
TELECOMMUNICATION INFORMATION:
TELEFRONE: 313 996-7295
TELEFRAX: 313 996-1553
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/08373911
Patent No. 5663296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 313 996-1553
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                              TYPE: 5 amino acids
TYPE: amino acids
TOPOLOGY: 1:--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                     , MOLECULE TYPE: peptide US-08-373-911-5
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4 IW 5
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US-08-373-911-6
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Patent No. 592529
GENERAL INFORMATION:
APPLICANT: COUGHLIN, ET AL.
TITLE OF INVENTION: METHOD FOR DISCOVERY OF PEPTIDE AGONISTS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOLEY GODWARD LLP
STREET: FIVE PALO ALTO SQUARE
STREET: FIVE PALO ALTO
CITY: PALO ALTO
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      METHOD FOR DISCOVERY OF PEPTIDE AGONISTS
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                                                                                  56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,506A
FILING DATE: 7-JUN-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UCAL-239/00US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94306-2155
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08483506A
Patent No. 525559
GENERAL INFORMATION:
APPLICANT: COUGHLIN, ET AL.
ITLE OF INVENTION: METHOD FOR DI
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: COOLEY GODWARD LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: NEELEY, RICHARD L. PhD
REGISTATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: UCAL
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (415) 843-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 857-0663
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                        Query Match
Best Local Similarity 50.0%
Matches 1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                           MOLECULE TYPE: peptide US-08-373-911-6
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Matches 1; Conserv
TYPE: amino acid
                   linear
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                 TOPOLOGY:
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                                                                                                                                                                                                                                                          RESULT 53
US-08-483-506A-1
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4 IW 5
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TELEPHONE:
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US-08437-607A-38
is equence 38, Application US/08437607A
j Patent No. 595579
j GENERAL INFORMATION:
j APPLICANT: Leonard, James N. Montminy, Marc R.
TITLE OF INVENTION: ISLET-SPECIFIC HOMEOPROTEIN AND TRANSCRIPTIONAL
j TITLE OF INVENTION: REGULATOR OF INSULIN GENE EXPRESSION
j VUMBER OF SEQUENCES: 49
j CORRESPONDENCE ADDRESS:
j ADDRESSEE: Klauber & Jackson
STREET: 411 Hackensack Avenue
j CITY: Hackensack
j CITY: Hackensack
j COUNTRY: USA
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                                                                                        COUNTRY: USA
ZIP: 94306-2155
COMPUTER READALLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,506A
FILING DATE: 7-JUN-1995
CLASSIFICATION NUMBER: US/08/483,506A
FILING DATE: 7-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: NEELEY, RICHARD L. PhD
REGISTRATION NUMBER: UCAL-239/00US
REFERENCE/DOCKET NUMBER: UCAL-239/00US
TELECOMMONICATION INFORMATION:
TELEPRAX: (415) 843-5000
TELEPAX: (415) 843-5000
TELEPAX: (415) 87-0663
; INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
TENNYTH: 5 amino acids
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OTHER INFORMATION: /note= "Should be Formyl-Met"
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/437,607A
FILING DATE: MAY 9, 1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
: FIVE PALO ALTO SQUARE
: 3000 EL CAMINO REAL
PALO ALTO
CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
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                                                               STATE: C
  STREET:
STREET:
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; Patent No. 6127339
; OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
; OTHER INFORMATION: synthesizing system (RaMPS)
US-08-981-122-2
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OTHER INFORMATION: Sequence of a peptide synthesized in Example 1 from L-form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: P7-176904
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR PLING DATE: 1996-06-21
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FILING DATE: 1996-06-21
SUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN VET: 2.0
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APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Masaharu
ITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
DESCRIPTION: amino acids 47-51 of the homeobox homeodomain
                                                                                                                                                                                                                                                                                                                                                          56.5%; Score 13; DB 2; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          56.5%; Score 13; DB 3; Length 5; llarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
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CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: JP 7-176904
PRIOR FILING DATE: 1995-06-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-981-122-19
; Sequence 19, Application US/08981122B
; Patent No. 6127339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 2, Application US/08981122B; Patent No. 6127339; GENERAL INFORMATION:
                  TELEFAX: 201 343-1684
TELEX: 133521
INFORMATION FOR SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
201 487-5800
                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                            5 amino acids
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Matches 1; Conserv
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                                                                                                                                                                                                                                                                                       ORGANISM: Rat
                                                                                                                                                                                                                                                            ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                            US-08-437-607A-38
                                                                                                                         LENGIH:
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LOCATION: 5
OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for
                                                                                                                                                                                                                                                                                                                                                              Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid
OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Sequence of a peptide synthesized in Comparative Example 2 from
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OTHER INFORMATION: L-form F-moc amino acids by solid phase method using a CTHER INFORMATION: multipeptide synthesizing system (RAMPS)
US-08-981-122-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
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CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: UP 7-176904
PRIOR FILING DATE: 1995-06-21
PRIOR FILING DATE: 1996-06-21
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver: 2.0
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               PRIOR APPLICATION NUMBER: JP 7-176904
PRIOR FILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 72, Application US/08981122B; Patent No. 6127339; GENERAL INFORMATION:
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Patent No. 6222021
GENERAL INFORMATION:
CURRENT FILING DATE: 1997-12-18
                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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Best Local Similarity
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US-08-981-122-72
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LOCATION: 5

COTATION: 5

OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for Patent No. 6127339

OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid US-08-981-122-37
                                                                                                                                                                                                                                          Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
OTHER INFORMATION: synthesizing system (RaMPS)
                                                                                                                                                                                                                        OTHER INFORMATION: Sequence of a peptide synthesized in Example 2 from L-form
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APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
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APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REPERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                   56.5%; Score 13; DB 3; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: JP 7-176504
PRIOR FILING DATE: 1995-06-21
PRIOR FILING DATE: 1996-06-21
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
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PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FLING DAFE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PAtentIN Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 37, Application US/08981122B; Patent No. 6127339; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 53, Application US/08981122B
; Patent No. 6127339
                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity 50.09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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                                                                                               SEQ ID NO 19
                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                              LENGTH
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CURRENT FILING DATE: 1998-12-18
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Matches 1; Conservative
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                                                                                                                                                                                                                        NAME/KEY: VARIANT
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US-09-638-202A-51
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1W 5
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Patent No. 6268328
GENERAL INFORMATION:
APPLICANT: Mitchinson, Colin
APPLICANT: Wendt, Dan J.
TITLE OF INVENTION: No. 6268328el Variant EGIII-Like Cellulase Compositions
FILE REFERENCE: GC555
CURRENT APPLICATION NUMBER: US/09/216,295
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TITLE OF INVENTION: Endotoxin Binding and Neutralizing TITLE OF INVENTION: Protein and Uses Thereof NUMBER OF SEQUENCES: 4 CORRESPONDENCE ADDRESS: ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Avenue, N.W., Suite 600 CITY: Washington
                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: United States of America
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/871,600A
FILING DATE: 30-AUG-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/704,872
FILING DATE: 21-JUN-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/883,457
FILING DATE: 15-MAY-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/701,501
FILING DATE: 16-MAY-1991
APPLICATION NUMBER: US 07/701,501
FILING DATE: 16-FEB-1990
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/210,575
ATTAC DATE: US DATA:
APPLICATION NUMBER: US 07/210,575
ATTAC DATE: US DATA:
APPLICATION NUMBER: US 07/210,575
ATTAC
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NAME: Esmond Policy
                                                                                                                                                                        STATE: District of Columbia COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Esmond, Robert W. REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 14:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2500
TELEFAX: (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 248636 SSK
INFORMATION FOR SEQ ID NO: 4:
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TOPOLOGY: not relevant
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Best Local Similarity 50.v.
1, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide US-08-871-600A-4
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STREET: 121 South Eighth Street, Ste. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                 56.5%; Score 13; DB 4; Length 5; llarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
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TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
CORPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTESEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/638,202A
FILING DATE: 11-Aug-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/096,749
FILING DATE: CURROWN:
APPLICATION NUMBER: 09/096,749
FILING DATE: CURROWN:
ATTORNEY/AGNT INFORMATION:
NAMME: Ann S. VİKSHİNS
REĞISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034USI
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 26
LENGTH: 5
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                                                                                                                                                                                                                                                                         LOCATION: (1) ... (5) OTHER INFORMATION: Xaa = Leu, Phe, or Ile
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 51, Application US/09638202A
; Patent No. 6462189
; GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
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STRANDEDNESS: single
                                                                                                                       TYPE: PRT ORGANISM: Artificial Sequence
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MOLECULE TYPE: peptide
                                                                                                                                                                                                               OTHER INFORMATION: Peptide
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Matches 1; Conserva
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                                                          COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : 4
IW 5
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                                                                                                                                                         APPLICANT: Hirschberg, Carlos B.
APPLICANT: Orellana, Ariel
APPLICANT: Hashimoto, Yasuhiro
APPLICANT: Swiedler, Stuart J.
APPLICANT: Wei, Zheng
APPLICANT: Ishihara, Masayuki
TITLE OF INVENTION: GLYCOSAMINOGLYCAN SPECIFIC
TITLE OF INVENTION: SULFOTRANSFERASES
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Cody, Wayne L.
APPLICANT: Doherty, Annette M.
APPLICANT: Topliss, John G.
TITLE OF INVENTION: Endothelin Antagonists II
CORRESPONDENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/899,432
FILING DATE: 16 June 1992
ATTORNEY/AGENT INFORMATION:
NAME: CLARK, PAUL T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04020/015002
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: 3.5" Diskette, 1.44 Mb COMPUTER: 18M PS/2 Model 50Z or 55SX OPERATING SYSTEM: MS-DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1) APPLICATION DATA: APPLICATION NUMBER: PCT/US93/05412 FILING DATE: 19930607
                                                                                                                 Sequence 12, Application PC/TUS9305412
GENERAL INFORMATION:
APPLICANT: Hirschberg, Carlos B.
APPLICANT: Orellana, Ariel
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GENERAL INFORMATION:
APPLICANT: Cody, Wayne L.
                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (617) 542-8900
TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
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TELEFAX: (617) 542-8906
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                                                                                                                                                                                                                                                                                                                                                                                                        Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: N/H
TOPOLOGY: N/A
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2 MW 3
4 LW 5
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S6.5%; Score 13; DB 5; Length 5;
Best Local Similarity 50.0%; Pred. No. 2e+05;
Matches 1; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: KAUVAR, LAWRENCE M.
TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver. #1.25
CURRENT APPLICATION DATA:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                       NAME: TINDEY, FRANCIS J.
REGISTRATION NUMBER: 33069
REFERENCE/DOCKET NUMBER: PD-4334-P2
TELECHONE: 313 996-7295
TELEPHONE: 313 996-7295
INPORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
ADDRESSEE: Warner-Lambert Company STREET: 2800 Plymouth Rd.
                                                                                                                                                                                                                                                                                                                                            US 08/033,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 121
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 amino acids
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MOLECULE TYPE: peptide
PCT-US95-04171-9
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           STREET: Zav.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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Matches 1, Conserv
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APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Rafnar, Thorum
APPLICANT: Rafnar, Thorum
APPLICANT: Ruo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSER: LAHLYE & COCKFIELD
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 5;
                                                                                                                                                                                        CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/946,237

FILING DATE: 19920914

CLASSIFICATION: 435

ATTORNEY/AGENT INPORMATION:

NAME: Bavid E. Brock

REGISTRATION NUMBER: 22,592

REGISTRATION NUMBER: 22,592

REGISTRATION NUMBER: BTT92-01

TELECOMMUNICATION INPORMATION:

TELEPHONE: (617) 861-6240

TELEPHONE: (617) 861-6240

TELEPRORE (617) 861-6240

TELEPRORE (617) 861-9540

TELEPRORE (617) 861-9540

TELEPRORE (617) 861-9540

TELEPRORE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: August 15, 1994
PRIOR APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: March 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: 1M1-018CN
      Brook, Smith & Reynolds, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 12; DB 1;
Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Mismatches
                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 3, Application US/08290448A; Patent No. 5676954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 52.2%;
Local Similarity 50.0%;
    ADDRESSEE: Hamilton, Brood
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: AMINO ACID
MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 02109-1875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Boston
                                                                                COUNTRY: US
ZIP: 02173
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US-08-290-448A-3
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Best Local
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                                                                                                                                      56.5%; Score 13; DB 6; Length 5; 50.0%; Pred. No. 2e+05; ative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 5, Score 13; DB 6; Length 5; Best Local Similarity 50.0%; Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                                                          5217869-24
;Patent No. 5217869
; APPLICANT: KAUVAR, LAWRENCE M.
; TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABOULT
FATERI NO. 5217869
; APPLICANT: KAUVAR, LAWRENCE M.
; TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4, Application US/07946237
Patent No. 5348874
GENERAL INFORMATION:
APPLICANT: Savakis, Charalambos
APPLICANT: Franz, Gerald H
APPLICANT: Loukeris, Athanasios
TITLE OF INVENTION: Eukaryotic Transposable Element
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
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APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
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                                                                                                                  Query Match
Best Local Similarity
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5217869-34
                                                                              LENGTH: 5
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1 MW 2
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                                                            ;SEQ ID NO:22:
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                                                                                                5217869-22
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                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Rogers, Bruce
APPLICANT: Rogers, Bruce
APPLICANT: Rapper, David G.
APPLICANT: Rapper, David G.
APPLICANT: Ranar, Thorunn
APPLICANT: Ruo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSE: LAHIVE & COCKFIELD
STREET: 60 State Street, suite 510
CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                 52.2%; Score 12; DB 1; Length 5; ilarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: Aquist 15, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: March 17, 1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 39, Application US/08290448A Patent No. 5676954 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI
TELECOMMUNICATION: TELEPHONE: (617)227-7400
              TELEPHONE: (617)227-7400
TELEPAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENCTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity bu.v.
Local 1; Conservative
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                                                                                                                                              MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                         RESULT ,72
US-08-290-448A-39
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52.2%; Score 12; DB 1; Length 5; 20.0%; Pred. No. 2e+05; tive 1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert, ADDRESSEE: Attn: W.H. Dreger STREET: 4 Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/064,111C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/064,111C
FILING DATE: 12-AUG-1993
CLASSIPICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU91/00580
FILING DATE: 13-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PK9567
FILING DATE: 19-NOV-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PK3879
FILING DATE: 13-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: DYSOGNATION:
NAME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A-58456/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                        ; Sequence 22, Application US/08064111C; Patent No. 5688760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 3, Application US/08290448A; Patent No. 5698204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 52.2'
Best Local Similarity 20.0'
Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
US-08-064-111C-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-064-111C-22
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Gaps

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0; Indels

4 LW 5

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us-09-403-440a-1.closed.rai

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Job time : 15 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      us-08-290-448A-39

Sequence 39, Application US/08290448A

Sequence 39, Application US/08290448A

GENERAL INFORMATION:

APPLICANT: Rogers, Bruce

APPLICANT: Rafaper, David G.

APPLICANT: Rate Thourn

APPLICANT: Rate Thourn

APPLICANT: Rate Mei-chang

TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses

NUMBER OF SEQUENCES: 93

CORRESPONDENCE ADDRESS:

ADDRESSER: LAHIVE & COCKFIELD

STREET: 60 State Street, suite 510

CITY: Boston

STATE: Massachusetts

COUNTRY: USA

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                 APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Kuo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCES: ADDRESS:
ADDRESSEE: LAHITE & COCKFIELD
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             52.2%; Score 12; DB 1; Length 5; 50.0%; Pred. No. 2e+05; arive 1; Mismatches 0; Indels
                                                                                                                                                                                                         CITY: Boston
STATE: Massachusetts
COMPITEY: USA
ZIP: 02109-1875
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
SOFTWARE: PAPELICATION DATA:
APPLICATION NUMBER: US 08/290,448A
FRIGH APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: MAY 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: MAY 29, 1980
APPLICATION NUMBER: US 07/325,365
FILING DATE: MACCh 17, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 36,207
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI
TELECOMMUNICATION: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linear
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-290-448A-39
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             CURRENT APPLICATION DATA:
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: AUGUST 15, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: March 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
                                                                                                                                                                                                                                                                                        REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-018CN
TELEPONNUNICATION INFORMATION:
TELEPEAX: (617)227-7400
TELEPEAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGHH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Search completed: February 12, 2003, 10:53:11
PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid TOPOLOGY: linear
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-290-448A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:12:22 ; Search time 5.41667 Seconds (without alignments) 88.739 Million cell updates/sec Run on:

US-09-403-440A-1 23 Title: Perfect score:

1 PXXLW 5 Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched:

4984 Total number of hits satisfying chosen parameters:

283224 segs, 96134422 residues

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 73:\*
1: Dir1:\*
2: Dir2:\*
3: Dir3:\*
4: Dir4:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## STIMMARTES

		Iq H chain V-D-J r	homeotic protein G		GTP-binding protei	dnaA protein - Pse	T-cell receptor be	Ig H chain V-D-J r	glycine reductase	adipokinetic hormo	litorin - Rohde's	T-cell receptor ga	receptor		rhlR protein - Pse	phospholipase A2 (	gene HEXA protein	microbial collagen	thyroglobulin - ra	Ig H chain V-D-J r	Ig heavy chain C r	Ig heavy chain C r	collecting duct wa	glyceraldehyde-3-p	hypothetical prote	Iq H chain V-D-J r	Ig H chain V-D-J r	H chain V-D-J	Ig H chain V-D-J r	sell receptor
SUMMARIES	ID	PH1613	C37290	868897	S38281	B34835	PT0519	PH1602	A39308	A24244	S07241	E49033	F49033	LFTWWE	C53652	A29169	164829	A26093	158273	PH1620	G37266	D37267	151905	S54344	A35105	PH1625	PH1626	PH1627	PH1594	PH0801
	DB	2	~	7	~	7	7	~	~	7	7	~	7	٦	7	~	7	7	7	~	~	7	N	7	~	N	7	~	~	7
	Length	15	16	25	24	9	9	7	σ	6	6	10	10	11	11	12	12	12	12	13	13	13	13	13	14	14	14	14	14	14
	Query Match		3.9	3.9	9.6	2.5	65.2	2.5	2.5	5.2	5.2	65.2	5.2	2.5	2.5	5.2	5.2	2.5	5.2			٠		2.2					2.5	
٨	g g	87	73	73	Ö	ø	ø	ø	ø	9	ø	Ġ	ý	Ü	Ġ	Ö	Ġ	Ġ	Ġ	Ġ	Ğ	Ö					Ö			9
	Score	20	17	17	16	15	15	15	15	15	15	15	15	15	15	15	15		15				15	15	15	15	15	15	15	15
	Result No.	П	7	М	4	Ŋ	9	7	ω	σ	10	11	12	13	14	15	16	17	18	19	50	21	22	23	24	25	56	27	28	29

TcR delta chain V-	T cell receptor al	7 alpha-hydroxy-4-	Ig H chain V-D-J r	Ig H chain V-D-J r	TcR delta chain V-	hydrogenase (EC 1.	Ig H chain V-D-J r	Ig H chain V-D-J r	TCR delta chain V-	transforming prote	caldesmon - rabbit	Ig H chain V-D-J r	T-cell receptor de	shikimate 5-dehydr	choline O-acetyltr
F49037	PH1788	A45103	PH1590	PH1612	E49037	S03405	PH1637	PH1638	D49037	A46236	B44873	PH1630	A32220	140062	S19914
~	7	~	7	7	~	~	7	~	7	~	7	~	7	~	7
14	15	15	15	15	15	16	16	16	16	16	17	17	18	18	18
65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2	65.2
15	15	12	15	15	15	12	15	15	12	15	15	15	15	15	15
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

## ALIGNMENTS

PH1613

```
Ig H chain V-D-J region (clone B-less 17) - mouse (fragment)
(Species: Mus musculus (house mouse)
(Spacies: Mus musculus (house mouse)
(Spacies: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
(SAccession: PH1613
(R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
A;Levinson, D.A.; Campos-Torres, J.; Leder, P.
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                        A;Molecule type: DNA
A;Residues: 1-15 <LEV>
A;Experimental source: bone marrow pre-B lymphocyte
C;Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                       A; Accession: PH1613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Gaps .; 0 h Similarity 60.0%; Pred. No. 2.2e+02; 3; Conservative 0; Mismatches 2; Indels Best Loc Matches

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10 FTMLW 14 1 FXXLW 5 qq ò

RESULT 2

C37220 homeotic protein Gah-3 - mouse (fragment) c.Species: Mus musculus (house mouse) C.Species: Mus musculus (house mouse) C.Accession: C37290; C38809 C;Accession: C37290; C38809 C;Accession: C37290; C38809 C;Accession: C37290; C3880; D.S. A. 88, 10706-10710, 1991 A;Title: Identification of 10 murine homeobox genes. A;Title: Identification of 10 murine homeobox genes. A;Refatus: preliminary; not compared with conceptual translation A;Status: preliminary; not compared with conceptual translation A;Molecule type: DNA A;Molecule type: DNA A;Residues: 1-16 <SIN> C;Superfamily: unassigned homeobox proteins; homeobox homology C;Keywords: DNA binding; homeobox, nucleus; transcription regulation F;1-14/Domain: homeobox homology (fragment) <ADOX>

Gaps .. 0 Query Match 73.9%; Score 17; DB 2; Length 16; Best Local Similarity 40.0%; Pred. No. 1e+03; Matches 2; Conservative 1; Mismatches 2; Indels

.; 0

12 FKKMW 16 1 FXXLW 5 g ò

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CiSpecies: Mus musculus (house mouse)
CiSpecies: Mus musculus (house mouse)
CiDate: OS-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
CiAccession: PH1602
Ribevinson, D.A.; Campos-Torres, J.; Leder, P.
Ribevinson, D.A.; Campos-Torres, J.; Leder, P.
A.Title: Modecular characterization of transgene-induced immunodeficiency in B-less mice
A.Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       glycine reductase (EC 1.4.99.-) sulfhydryl protein C, alpha chain - Clostridium sticklan
C,Species: Clostridium sticklandii
C,Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 15-Aug-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A,Description: glycine reductase complex catalyzes the reductive deamination of glycine C;Keywords: ATP; oxidoreductase
                                                                                                                                                                                                                                                                                                                                                                                                R.Feeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A.Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A.Reference number: PT0509; MUID:91277601; PMID:1711558
A.Recession: PT0519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                T-cell receptor beta chain V-D-J region (100-4C) - mouse (fragment) C;Species: Mus musculus (house mouse) C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997 C;Accession: PTO519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ig H chain V-D-J region (wild-type clone 313) - mouse (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         65.2%; Score 15; DB 2; Length 6; 100.0%; Pred. No. 2.8e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.2%; Score 15; DB 2; Length 7; 100.0%; Pred. No. 2.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Experimental source: bone marrow pre-B lymphocyte C; Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A, Experimental source: adult thymus, strain BALB/c
C, Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 100.0%; Pred. No. 2.8
Matches 2; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Status: translation not shown A;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Matches 2; Conserva
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A;Molecule type: protein
A;Residues: 1-8 <STA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Residues: 1-6 <FEE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: DNA A; Residues: 1-7 < LEV>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Accession: A39308
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4 LW 5
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                                                                     5 LW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A39308
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R;Yee, T.W.; Smith, D.W.
R;Yee, T.W.; Smith, D.W.
A;Title: Pseudomonas chromosomal replication origins: a bacterial class distinct from Estable and the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the control of the
                                                                                                                                                                           C;Accession: S68897
S;Gordon, R.D.; Leighton, I.A.; Campbell, D.G.; Cohen, P.; Creaney, A.; Wilton, D.C.; Ma Eur. J. Biochem. 238, 690-697, 1996
A;Title: Cloning and expression of cytosolic phospholipase A(2) (cPLA(2)) and a naturall actural in an increase in specific activity.
A;Reference number: S68897; MUID:96300233; PMID:8706669
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A;Mesidues: 1-24 eMOO>
C;Superfamily: ras transforming protein; translation elongation factor Tu homology
C;Keywords: GTP binding
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                                                                  phospholipase A2 (BC 3.1.1.4), cytosolic - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 04-Dec_1997 #sequence_revision 12-Dec-1997 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTP-binding protein - clawed frog (fragments)
C;Species: Xenopus sp. (clawed frog)
C;Date: 19-May-1994 #sequence_revision 27-Feb-1997 #text_change 28-May-1999
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C,Date: 13-Jul-1990 #sequence_revision 13-Jul-1990 #text_change 08-Oct-1999
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A;Residues: 1-6 <YEE>
A;Cross-references: GB:M30125; NID:g151419; PIDN:AAA25916.1; PID:g151421,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      73.9%; Score 17; DB 2; Length 25; 40.0%; Pred. No. 1.5e+03; Live 1; Mismatches 2; Indels
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A;Residues: 1-25 <GOR>
C;Keywords: carboxylic ester hydrolase; cytosol
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probable trpEG leader peptide - Thermus aquaticus Cispecies: Thermus aquaticus Cispecies: Thermus aquaticus Cispecies: Thermus aquaticus Cibate: 30-Jun-1991 #sequence_revision 30-Jun-1991 #text_change 16-Jul-1999 CiAccession: 803315 Risato, S.; Nakada, Y.; Kanaya, S.; Tanaka, T. Risato, S.; Nakada, Y.; Kanaya, S.; Tanaka, T. Aititle: Molecular Acta 950, 303-312, 1988 Aititle: Molecular cloning and nuclectide sequence of Thermus thermophilus HB8 trpE and A;Reference number: 803315; MUID:89000781; PMID:2844259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence_revision 17-Mar-2000 #text_change 17-Mar-2000
C;Accession: F49033 #sequence_revision 17-Mar-2000 #text_change 17-Mar-2000
C;Accession: F49033 #sequence_revision 17-Martinez, C.; Spits, H.; Brenner, M.B. Bur. J. Immunol. 21, 2990-3007, 1991
A;Title: Functionally distinct subsets of human gamma/delta T cells.
A;Reference number: A49033; MUID:92083926; PMID:1684157
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A,Residues: 1-11 <SAT>
A,Cross-references: EMBL:X07744; NID:g48261; PIDN:CAA30565.1; PID:g48262
A,Note: the source is designated as Thermus thermophilus HB8
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A;Note: sequence extracted from NCBI backbone (NCBIN:72605, NCBIP:72606)
C;Keywords: T-cell receptor
                                                                                                                                                                        A;Cross-references: GB:S72587; NID:g240696; PIDN:AAB20630.1; PID:g240697 A;Note: sequence extracted from NCBI backbone (NCBIN:72591, NCBIP:72595) A;Accession: D49033
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A;Residues: 1-10 <MOR2>
A;Cross-references: GB:S72587; NID:g240696; PIDN:AAB20630.1; PID:g240697
A;Note: sequence extracted from NCBI backbone (NCBIN:72587, NCBIP:72589)
C;Keywords: T-cell receptor
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A; Title: Functionally distinct subsets of human gamma/delta T cells. A; Reference number: A49033; MUID:92083926; PMID:1684157 A; Accession: E49033 A; Status: preliminary
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A;Molecule type: DNA
A;Residues: 1-10 <MOR>
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A;Residues: 1-9 <JAF>
(S;Superfamily: adipokinetic hormone
C;Reywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Gly) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Date: 31-Mar-1998 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
C;Accession. A24244
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Holman, G.M.; Wagner, R.M.; Ridgway Biochem. Biophys. Res. Commun. 135, 622-628, 1986
A;Title: Isolation and primary structure of a peptide from the corpora cardiaca of Helic A;Reference number: A24244; MUID:86186794; PMID:3964263
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R;Barra, D.; Falconieri Erspamer, G.; Simmaco, M.; Bossa, F.; Melchiorri, P.; Erspamer, R;Barra, D.; Falconieri Erspamer, G.; Simmaco, M.; Bossa, F.; Melchiorri, P.; Erspamer, A;Title: Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei.
A;Reference number: S07241; MUID:85127560; PMID:3838283
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C;Keywords: amidated carboxyl end; blocked amino end; neuropeptide; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Met) #status experimental
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence_revision 25-Aug-1995 #text_change 21-Jul-2000
C;Accession: E49033; h94033
R;Morita, C.T.; Verma, S.; Aparicio, P.; Martinez, C.; Spits, H.; Brenner, M.B. Bur. J. Immunol. 21, 2999-3007, 1991
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C;Species: Phyllomedusa rohdei (Rohde's leaf frog)
C;Date: 12-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
                                                                                                                                                                                                                                                                                                                                                                                                    N.Alternate names: Hez-AKH
C;Species: Hellothis zea (bollworm, corn earworm, tomato fruitworm)
C;Date: 31-Mar-1988 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
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                                                                                                     0; Indels
                                               Length 8;
                                         65.2%; Score 15; DB 2; Le
100.0%; Pred. No. 2.8e+05;
tive 0; Mismatches 0;
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                                                                                                         2; Conservative
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                                                                            Best Local Similarity
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0; Mismatches

2; Conservative

Matches

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C;Species: Pseudomonas aeruginosa
C;Date: 21-011-1995 #sequence_revision 28-Jul-1995 #text_change 21-Aug-1998
C;Accession: C53652
R;Ochsner, U.A.; Fiechter, A.; Reiser, J.
Biol. Chem. 269, 19787-19795, 1994
A;Title: Isolation, characterization, and expression in Escherichia coli of the Pseudomo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            phospholipase A2 (EC 3.1.1.4) precursor - sheep (fragment)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Adcession: A29169
R;Dutilh, C.E.; Van Doren, P.J.; Verheul, F.E.A.M.; De Haas, G.H.
Bur. J. Biochem. 53, 91-97, 1975
A;Tile: Isolation and properties of prophospholipase A2 from ox and sheep pancreas.
A;Reference number: A94661
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A;Residues: 1-12 <DUT>
C;Superfamily: phospholipase A2
C;Keywords: carboxylic ester hydrolase; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: DNA
A;Residues: 1-11 <OCH>
A;Cross-references: GB:L28170
C;Superfamily: sdiA regulatory protein
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A; Status: preliminary
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MEDLINE=94019818; PubMed=8413630;
Moore M.S., Blobel G.;
"The GTP-binding protein Ran/TC4 is required for protein import into
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                    Wabnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C., Bowie J.H., Tyler M.J., Wallace J.C., Interesting tree frog Litori electrica. Comparison with the skin peptides from Litoria rubella.";
                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
VCBI_TaxID=104895;
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Amphibia, Batrachia, Anura, Mesobatrachia, Pipoidea, Pipidae,
Xenopodinae, Xenopus.
VCBI_TaxID=8355,
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         P27503
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6 AA; 792 MW; 6683704772C9A000 CRC64;
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01-OCT-1996 (Rel. 34, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
GTP-binding nuclear protein RAN (TC4) (Fragments).
                                                                                                                                                                                                      15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Electrin 1.
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                          CCAA_STRII
LPCP_YEAST
Y15_BPT3
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HTF1_PERAM
HTF2_PERAM
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          PQQA_KLEPN
XYC1_ACIGB
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-!- SUBCELLULAR LOCATION: SECRETED.
Amphibian skin; Amidation.
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TISSUE=Skin secretion;
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AC P82096;
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                                                              February 12, 2003, 11:11:31; Search time 2.70833 Seconds (without alignments) 76.572 Million cell updates/sec
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       GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                        112892 segs, 41476328 residues
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A1BG EQUAS
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RAN XENLA
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Gapop 10.0 , Gapext 0.5
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         -!- FUNCTION: GTP-BINDING PROTEIN INVOLVED IN NUCLEOCYTOPLASMIC TRANSPORT. REQUIRED FOR THE IMPORT OF PROTEIN INTO THE NUCLEUS AND ALSO FOR RNA EXPORT. INVOLVED IN CHROMATIN CONDENSATION AND CONTROL OF CELL CYCLE (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- SUBCELLULAR: BELONGS TO THE RAN FAMILY IN THE RAS SUPERFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei."; FBBS Lett. 188:53-56(1985).
-i SUBGELLUAR: LOCATION: Secreted.
-i SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                           Phyllomedusa rohdei (Rohde's leaf frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Phyllomedusinae; Phyllomedusa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-85127560; PubMed-3838283;
Barra D., Erspamer G.F., Simmaco M., Bossa F., Melchiorri P.,
Erspamer V.;
                                                                                                                                                                    Score 16; DB 1; Length 24;
Pred. No. 9.18+02;
Transhas 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
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                                                                                                                                                         24 AA; 2657 MW; B69F83236247A250 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 AA; 1090 MW; 4ECCCIE861ADC377 CRC64;
                                                                                                                                      GTP (BY SIMILARITY).
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                                                                        HSSP, P17080; 1A2K.
InterPro; IPR002041; RAN.
PROSTITE; PS01115; RAN; PARTIAL.
GTP-binding; Nuclear protein; Protein transport.
                                                                                                                                                                                                                                                                                                                   01-NOV-1988 (Rel. 09, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P80474;
01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
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Conservative 0; Mismatches
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InterPro; IPR000074; Bombesin.
Pfam; PP02044; Bombesin; 1.
PROSITE; ES00257; BOMBESIN; 1.
Bombesin family; Amidation.
                                                                                                                                                                               69.6%;
                                                                                                                                                                             Query Match
Best Local Similarity 40.0°
Matches 2; Conservative
Nature 365:661-663(1993).
                                                                                                                                                                                                                                                                                               STANDARD;
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nes 2; Conserv
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P08946;
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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Biochim. Biophys. Acta 950:303-312(1988).
-!- FUNCTION: THIS PROTEIN IS INVOLVED IN CONTROL OF THE BIOSYNTHESIS
                                                                                                                                                                                                                                                                                                                                                                                                                       Microbiology 141:3087-3093(1995).

-!- FUNCTION: AMINOPEPTIDASE WHICH HYDROLYZES SUBSTRATES WITH FREE N-FERMINAL AMINO ACID RESIDUES BUT NOT N-TERMINAL BLOCKED ONES.

OPTIMUM ACTIVITY IS MEASURED AT PH 7.5. MAY BE IMPORTANT IN THE NUTRITION AND PATHOGENESIS OF THE ORGANISM IN THE HUMAN ORAL
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MEDLINE=88900781; PubMed=2844259;
Sato S., Nakada Y., Kanaya S., Tanaka T.;
"Molecular cloning and nucleotide sequence of Thermus thermophilus
                                                                                                                                                                                                                                                                                                                              Spratt D.A., Greenman J., Schaffer A.G.; "Capnocytophaga gingivalis aminopeptidase: a potential virulence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .;
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Bacteria; Thermus/Deinococcus group; Deinococci; Thermales;
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01-OCT-1996 (Rel. 34, Last annotation update)
Aminopeptidase (BC 3.4.11.-) (Fragment).
Capnocytophaga gingivalis.
Bacteria, Bacteroidetes; Flavobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 65.2%; Score 15; DB 1; Length 10; Local Similarity 100.0%; Pred. No. 6.5e+02; les 2; Conservative 0; Mismatches 0; Indels
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SEQUENCE 11 AA; 1228 MW; 364B295A772DC5A7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 10 10 10 10 AA; 1306 MW; 00C0A6DB43772694 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- COFACTOR: REQUIRES MAGNESIUM OR CALCIUM.
Hydrolase; Aminopeptidase; Magnesium; Calcium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1988 (Rel. 09, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch).
                                                                                                                                      Flavobacteriaceae; Capnocytophaga.
                                                                                                                                                                                                                                                           STRAIN=ATCC 33624;
MEDLINE=96118234; PubMed=8574402;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1988 (Rel. 09, Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Irp operon leader peptide.
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NCBI_TaxID=274;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; S03315; LFTWWE.
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Best Local Similarity
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CR31 LITSP
P56238;
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SEQUENCE
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16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Peptide hormone 4 (Pea-YLS-amide).
Eukarjota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Perrygota; Neopera; Orthopteroidea; Dictyoptera; Blattaria;
NCBI_TaxID=6978;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Abdominal perisympathetic organs;
MEDINE=20140865; PubMed=10674456;
Predel R., Eckert M., Hollman G.M.;
"The unique neuropeptide pattern in abdominal perisympathetic organs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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    Gaps
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PREDINES=13130579; PubMed=1868686;
Patterson S.D., Bell K., Shaw D.C.;
"Donkey and horse alpha 1 B-glycoprotein: partial characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Būkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Perissodactyla; Equidae; Equus.
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-!- FUNCTION: PLASMA GLYCOPROTEIN OF UNKNOWN FUNCTION.
-!- SUBCELLULAR LOCATION: Extracellular.
-!- SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY.
Immunoglobulin domain; Glycoprotein; Plasma.
NON TER.
20 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; rative 0; Mismatches 0; Indels
    Indels
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17 AA; 1968 MW; CAAFB57ECD8218A9 CRC64;
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-!- FUNCTION: UNKNOWN.
-!- MASS SPECTROMETRY: MW=1966.48; METHOD=MALDI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-FEB-1995 (Rel. 31, Created)
01-FRB-1995 (Rel. 31, Last sequence update)
11-FEB-1995 (Rel. 31, Last annotation update)
Alpha-18-glycoprotein (Postalbumin) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AA
    0; Mismatches
                                                                                                                                                                                    PRT;
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                                                                                                                                                                                                                            16-OCT-2001 (Rel. 40, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neuropeptide; Amidation.
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  Conservative
                                                                                                                                                                                    STANDARD;
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Best Local Similarity
Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of insects.";
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P39090;
Matches / 2;
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SEQUENCE
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characterization.";
Arch. Biochem. Biophys. 323:187-194(1995).
- PUNCTION: MAJOR ROLE IN THE SYNTHESIS OF NUCLEOSIDE TRIPHOSPHATES
- PUNCTION: MAJOR ROLE INILARITY).
- CATHER THAN ATP (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                 16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Nucleoside diphosphate kinase (EC 2.7.4.6) (NDFK) (NDP kinase) (NDPK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Litoria splendida (Magnificent tree frog), and
Litoria gilleni (Centralian tree frog).
Usukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIINE=96019963; PubMed=7487065;
Biondi R.M., Veron M., Walz K., Passeron S.;
"Candida albicans nucleoside-diphosphate kinase: purification and
                                                                                                                                                                                                                                              Candida albicans (Yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes;
Saccharomycetales, mitosporic Saccharomycetales; Candida.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE, AND MASS SPECTROMETRY.
SPECIES-L.splendida; TISSUE-Parotoid gland;
Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.;
Peptides from Australian frogs. Structures of the caerins and caeridin 1 from Litoria splendida.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21 AA; 2379 MW; 9DABB3A325947001 CRC64;
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15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleoside triphosphate.
--- SUBUNIT: HOMOHEXAMER.
--- SUBCELLULAR LOCATION: Cytoplasmic.
--- SIMILARITY: BELONGS TO THE NDK FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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PROSITE; PS00469; NDP_KINASES; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transferase; Kinase; ATP-binding.
                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE, AND CHARACTERIZATION
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InterPro; IPR001564; NDK.
Pfam; PF00334; NDK; 1.
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STANDARD;
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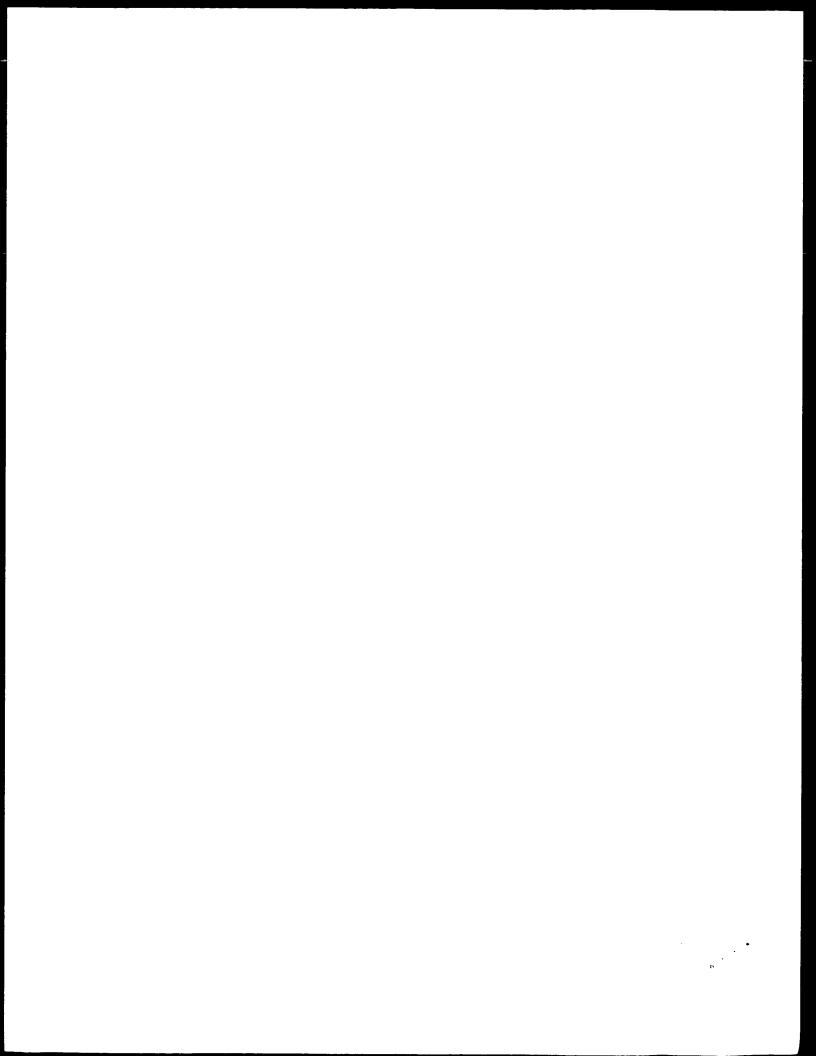
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Caerin 3.3.
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CR41_LITCE
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                               SPECIES-L.gilleni, TISSUE-Parottid gland;
Waugh R.J., Stone D.J.M., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structures of the caerins and caeridins from Litoria gilleni.";
J. Chem. Res. 139:937-961(1993).
-!- FUNCTION: ANTIBACTERIAL PREPIDE, THAT ADOPTS AN ALPHA HELLCAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERRANT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    J. Chem. Res. 138;910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL
CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structures of the caerins from
                                                                                                                                                                                                                                                                                                                                                                                                                      Medaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;0
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0
                                                                                                                                                                                                         65.2%; Score 15; DB 1; Length 22; llarity 100.0%; Pred. No. 1.4e+03; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.2%; Score 15; DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                   22 AA; 2385 MW; 1D4411E2E9D43739 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 AA; 2400 MW; 1D440B3829D4367C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.4e+03;
Chem. Soc. Perkin Trans. 1:3173-3178(1992).
                                                                                                                                                                                                                                                                                                                                                            15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Rel. 36, Created)
(Rel. 36, Last sequence update)
(Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                        22 AA.
                                                                                                                                                 -1- MASS SPECTROMETRY: MW=2382; METHOD=FAB.
Antibiotic; Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -1- MASS SPECTROMETRY: MW=2397; METHOD=FAB.
Antibiotic; Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ilarity 100.0%; Pred. No. 1.4
Conservative 0; Mismatches
                                                                                                                                                                          AMIDATION.
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                                                                                                                                                                                                                                                                                                                                                                                                          Litoria caerulea (Green tree frog)
                      SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                       STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Parotoid gland;
                                                                                                                                                                                                                    Local Similarity
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les 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI TaxID=30344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Litoria caerula."
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16-OCT-2001
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P56240}
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SEQUENCE
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CR32 LITCE
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CR33_LITCE
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                                                                                                                                                                                                                                                                                Litoria caerula.";
J. Chem. Res. 138:910-936(1993)
J. Chem. Res. 138:910-936(1993)
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL
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-!- FUNCTION: ANTIBACTERIAL PERTIDE, THAT ADOPTS AN ALPHA HELLICAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. BACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Parotoid gland;
Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structures of the caerins from
                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
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22 AA; 2427 MW; 1D440B2200D4367C CRC64;
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22 AA; 2455 MW; 3AB40B2200D43663 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 1; Le larity 100.0%; Pred. No. 1.4e+03; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-1998 (Rel. 36, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -1- MASS SPECTROMETRY: MW=2424; METHOD=FAB.
Antibiotic; Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GLANDS.
-!- MASS SPECTROMETRY: MW=2452; METHOD=FAB.
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Litoria caerulea (Green tree frog).
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                                                                                                                                                                         SEQUENCE, AND MASS SPECTROMETRY.
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                                                                                    Pelodryadinae; Litoria.
                                                                                                                                                                                                 TISSUE=Parotoid gland;
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tes 2; Conserv
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Best Local Similarity
                                                                                                                   NCBI TaxID=30344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=30344;
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SEQUENCE
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GLANDS
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SEQUENCE
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  2 LW
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CR43_LITCE
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-!- FUNCTION: ANTIBACTERIAL PREPIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION HICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFREENT ANTIMICROBIAL, SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
                                                                                                                                                                                                                      Litoria caerula.";
J. Chem. Res. 138:910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL
CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN
DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL
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                                                                                                                                                                                            Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structures of the caerins from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia; Batrachia, Anura; Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                Litoria caerulea (Green tree frog).
Skaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
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                                                                                                                                                                                                                                                                                                                                                                                       65.2%; Score 15; DB 1; Length 23; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                             23 AA; 2329 MW; 83BFDD80B6ADDC87 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P56243;
15-VUL-1998 (Rel. 36, Created)
15-VUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                         15-UTL-1998 (Rel. 36, Created)
15-UTL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23 AA.
  23 AA.
                                                                                                                                                                                                                                                                                                                  -!- MASS SPECTROMETRY: MW=2326; METHOD=FAB. Antibiotic; Amphibian skin; Amidation.
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 PRT;
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                                                                                                                                                                 SEQUENCE, AND MASS SPECTROMETRY. TISSUE=Parotoid gland;
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Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    2; Conservative
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  STANDARD;
                                                                                                                                                                                                                                                                                                                                                  23
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                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                     NCBI_TaxID=30344;
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CR41 LITCE
P56242;
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                                                                      Caerin 4.1
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Litoria caerula.";
J. Chem. Res. 138:910-936(1993).

-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION WHICH AND DISPURE BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
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                                                                                                                                                                                                             Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
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23 AA; 2356 MW; B98FDD80B6ADDC9D CRC64;
                                                         15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
   23 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -!- MASS SPECTROMETRY: MW=2353; METHOD=FAB. Antibiotic; Amphibian skin; Amidation.
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                                                                                                                                                                                      Litoria caerulea (Green tree frog)
                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE, AND MASS SPECTROMETRY.
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   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Parotoid gland;
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les 2; Conservat
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CR43 LITCE
P56244;
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February 12, 2003, 11:12:02; Search time 9.79167 Seconds (without alignments) 105.216 Million cell updates/sec
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                      OM protein - protein search, using sw model
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9297 Total number of hits satisfying chosen parameters: 671580 segs, 206047115 residues BLOSUM62 Gapop 10.0 , Gapext 0.5 US-09-403-440A-1 23 1 FXXLW 5 seq length: 0 seq length: 25 Title: Perfect score: Scoring table: Minimum DB Maximum DB Sequence: Searched:

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

SPTREMBL 21:\*

1: sp\_archea:\*
2: sp\_bacteria:\*
3: sp\_fungi:\*
4: sp\_invartebrate:\*
5: sp\_invartebrate:\*
5: sp\_invartebrate:\*
5: sp\_invartebrate:\*
5: sp\_pammal:\*
sp\_unclassified:\* sp\_vertebrate:\* sp\_bacteriap: \* sp\_rodent:\*
sp\_virus:\* archeap:\* rvirus:\* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Q8uvw3 brachydanio Q61943 mus musculu Q9byms homo sapien Q53579 rhodobacter Q53580 rhodobacter Ogdw76 mus sp. hom O35353 rattus norv Q66548 human herpe Q72017 human immun P72429 salmonella O85406 coxiella bu O95953 homo sapien Q38366 bacteriopha Q9w6d7 gallus gall Q9s0u6 shigella so Q9np68 homo sapien Description SUMMARIES Q9QW76 Q35353 Q66548 Q72017 Q9S0U6 P72429 Q8UVW3 Q61943 Q9BYM6 Q53579 Q53580 Q9W6D7 095953 Q38366 085406 Query Match Length DB 87.0 87.0 887.0 738.3 733.9 773.9 773.9 773.9 669.6 65.2 65.2 65.2 Score Š.

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## ALIGNMENTS

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SECUENCE FROM N.A.
MEDLINE=99067015; PubMed=9851690;
Grechez-Cassiau A., Bernard M., Ladjali K., Rodriguez I.R., Voisin P.;
"Structural analysis of the chicken hydroxyindole-O-methyltransferase
                                       01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOX-1999 (TrEMBLrel. 20, Last sequence update)
Hydroxyindole-o-methyltrensferase annotation update)
Hydroxyindole-o-methyltrensferase isoform A (Fragment).
Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianiae;
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                                                                                                                                                                                                                                                                                                                                                                 87.0%; Score 20; DB 13; Length 22; 60.0%; Pred. No. 5.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                       22 AA; 2783 MW; 35726EABF1E450C8 CRC64;
              22 AA
              PRT;
                                                                                                                                                                                                                                                           gene.";
Bur. J. Blochem. 258:44-52(1998).
EMBL; AF116455; AAD2444.1; -
Methyltransferase; Transferase.
            PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 60.0
les 3; Conservative
                                                                                                                                                   Gallus.
NCBI_TaxID=9031;
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ID Q9
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25 AA.

01-OCT-2000 (TrEMBLrel. 15, Created)

PRELIMINARY;

Q9NP68 Q9NP68;

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O9BYM6
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Probom; PD000500; Ion_glu_receptor.
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Actinopterygii, Neopterygii, Teleostei, Ostariophysi, Cypriniformes,
Cyprinidae, Danio.
NCBI_TaxID=7955,
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                                                                                                                                          Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                              A Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
A Joyce K.A., Salisbury J., Sibson D.R., Vergote I., Walker C.;
Submitted (NOV-1999) to the EMBL/GenBank/DDBJ databases.
R EMBL, AF209136; AAF36352.1; -.
R EMBL, AF209131; AAF36355.1; -.
R EMBL, AF209131; AAF36356.1; -.
R EMBL, AF209131; AAF36359.1; -.
R EMBL, AF209133; AAF36350.1; -.
R EMBL, AF209134; AAF36350.1; -.
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NON TER 25 25
SEQÜENCE 25 AA; 2837 MW; 9EB8AA36B50DB17E CRC64;
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01-WAR-2002 (TrEMBLrel. 20, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
AMPA receptor subunit 2 (Fragment)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update) 01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
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                                                         Tumor suppressor p53 (Fragment).
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Pfam, PF00870; P53; 1.
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Matches 3; Conservative
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Neuroscience 69:807-819(1995).
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=96164254; PubMed=8596650;
Bessis A., Salmon A.M., Zoli M., Le Novere N., Picciotto M.,
Changeux J.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      82.6%; Score 19; DB 11; Length 21; 60.0%; Pred. No. 9.3e+02; tive 0; Mismatches 2; Indels
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Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; X82655; CAA57978.1; -.
MGD; MGI:87891; Chrnb2.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-2001 (TrEMBLrel. 19, Last annotation update)
Nicotinic acetylcholine receptor beta 2-subunit protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Receptor.
NON TER 21 21
SEQÜENCE 21 AA; 2351 MW; 28E57E2BC2BC67FA CRC64;
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
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PRELIMINARY;
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Matches 2; Conserva
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SEQUENCE
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Bacteria; Proteobacteria; alpha subdivision; Rhodobacter group;
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                                                                                                                                                                  Rhodobacter capsulatus (Rhodopseudomonas capsulata).
Bacteria; Proteobacteria; alpha subdivision; Rhodobacter group;
Rhodobacter.
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MEDLINE=92234963; PubMed=1569029;

Richter P., Brand M., Drews G.;

"Characterization of LHI- and LHI+ Rhodobacter capsulatus pufA
                                                                                                                                                                                                                           SEQUENCE FROM N.A.
MEDILTE=22134963; bubmed=1569029;
Richter P., Brand M., Drews G.;
"Characterization of LHI- and LHI+ Rhodobacter capsulatus pufA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          h Similarity 40.0%; Pred. No. 1.8e+03; Length 15; 2; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                             73.9%; Score 17; DB 2; Length 12; nilarity 40.0%; Pred. No. 1.5e+03; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Light-harvesting complex I alpha polypeptide (Fragment).
                                                                                                        01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 19, Last annotation update)
Light-harvesting complex I alpha polypeptide (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE 15 AA; 2054 MW; 3561FE413591D31A CRC64;
                                                                                                                                                                                                                                                                                                                     12 AA; 1627 MW; 0F92F6EA8A70532B CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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EMBL; S97552; AAC60406.1; -.
NON TER 15
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J. Bacteriol. 174:3030-3041(1992).
EMBL; S97551; AAC60405.1; -
NON TER 12 12 0F992
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Matches 2; Conserv
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Neuroscience 85.475.486(1998).

In heuroscience 85.475.486(1998).

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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Rattus.
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STRAIN=SPRAGUE-DAWLEY: TISSUE=BRAIN;
MEDLINE=98281588; PubMed=9622245;
MEDLINE=98281588; M.A. Frandes K.J., Cockett M.I.;
"Distribution of heterotrimeric G-protein beta and gamma subunits in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Guanine nucleotide-binding protein beta subunit 4 (Transducin beta
                                                                                                                                                                                                                                    Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
NCBI_TaxID=10095,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73.9%; Score 17; DB 11; Length 16; 40.0%; Pred. No. 1.9e+03; ative 1; Mismatches 2; Indels
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                                                               01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAY-2002 (TrEMBLrel. 21, Last annotation update)
Homeobox GSH-3 (Fragment).
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MEDLINE=92073356; PubMed=1683707;
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PROSITE; PS50071; HOMEOBOX_2; 1.
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PRELIMINARY;
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PRELIMINARY;
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"The analysis of EBV proteins which are antigenic in vivo.";
Nucleic Acids Res. 16:2859-2872(1988).
BENBL; X07531; CAA30406.1; -.
                     / Match 73.9%; Score 17; DB 11; Length 20; Local Similarity 40.0%; Pred. No. 2.4e+03; nes 2; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 73.9%; Score 17; DB 12; Length 20; Best Local Similarity 40.0%; Pred. No. 2.4e+03; Matches 2; Conservative 1; Mismatches 2; Indels
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                                                                                                                                                                              01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
EBV B95-8 Cl(g) DNA with antigen-coding ORF (Fragment) thuman herpesvirus 4 (Eppetain-Barr virus).
Viruses, daDNA viruses, no RNA stage, Herpesviridae;
03mmaherpesvirinae; Lymphocryptovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Michael N.L.;
Submitted (APR-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; U24486; AAA79676.1; -.
 20 AA; 2242 MW; C143688295AE0BE0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 20 AA; 2576 MW; 05F877676B235F95 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
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MEDLINE=88217505; PubMed=2835748;
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Best Local Similarity 40.0%
Matches 2; Conservative
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11 FRFIW 15
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RESULT 12

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MEDLINE=20248365; PubMed=10789503;
Chida T., Okamura N., Ohtani K., Yoshida Y., Arakawa E., Watanabe H.;
"The complete DNA sequence of the O antigen gene region of Plesiomonas
shigelloides serotype 017 which is identical to Shigella sonnei form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Houng H.H., Venkatesan M.M., "Genetic analysis of Shigella sonnei form I antigen: identification of a novel IS630 as an essential element for the form I antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. SPECIES-S.enterica serovar Choleraesuis; STRAIN=RF-1;
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40.0%; Pred. No. 4.8e+03;
trive 0; Mismatches 3; Indels . 0; Gaps
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SPECIES=S.enteritidis, STRAIN=82139; PLASMID=VIRULENCE;
SPECIES=S.enteritidis, UbbMed=9099859;
Rodriguez-Pena J.M., Buisan M., Ibanez M., Rotger R.;
"Genetic map of the virulence plasmid of Salmonella enteritidis and nucleotide sequence of its replicons.";
Gene 188:53-61(1997).
                                                                                                                      Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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"50 kW virulence plasmid of Salmonella enterica serovar
Choleraesuis.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Appression.";
Microb. Pathog. 25:165-173(1998).
EMBL; AB028134; BAA85070.1; -.
NON TER 25
SEQUENCE 25 AA; 2726 MW; A6CC5D24211CDAAD CRC64;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                               I antigen.";
Microbiol. Immunol. 44:161-172(2000).
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Conservative
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                                                                                                                  SPECIESES. Typhimurium, STRAIN=LT2 / SGSC1412 / ATCC 700720;
PLASMID=PSIT;
MEDLINE21534948; PubMed=1167609;
MEDLINE21534948; PubMed=1167609;
MCClelland M., Sanderson K.E., Spieth J., Clifton S.W., Latreille P., Courtney L., Porwollik S., Ali J., Dante M., Du F., Hou S., Layman D., Leonard S., Mayven C., Scott K., Holmes A., Grewal N., Mulvaney E., Ryan B., Sun H., Florea L., Miller W., Stoneking T., Nhan M., Waterston R., Wilson R.K.;
"Complete genome sequence of Salmonella enterica serovar Typhimurium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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MEDLINE=08348442; PubMed=9683477;
MILLEMS H., Jaeger C., Baljer G.;
"Physical and genetic map of the obligate intracellular bacterium Coxiella burnetii.";
J. Bacteriol. 180:3816-3822(1998).
Bypothetical protein.
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Pred. No. 4.8e+03;
1; Mismatches 2; Indels
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Submiveed (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; U64797; AAB53038.1; -.
EMBL; AB040415; BAB20531.1; -.
EMBL; AE006471; AAL23447.1; -.
Plasmid; Complete proteonor
SEQUENCE 25 AA; 3008 MW; B21F596B6DEFABE7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-2002 (TrEMBLrel. 20, Last annotation update)
Galactocerebrosidase (EC 3.2.1.46) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Hypothetical 1.0 kDa protein (Fragment).
Coxiella burnetii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8 AA; 993 MW; 046B5AA453772727 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 413:852-856(2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
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                                                                                      SEQUENCE FROM N.A.
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085406
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095953
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OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; OC Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo. OX NCBL TaxID=9606; RN [1]
RP SEQUENCE FROM N.A.
RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Lulli L., Torchiana E., Finocchiaro G.; RA Glycosidase; Hydrolase.
FT NON TER 9 AA: 1069 MW; 374E2AADC2C699C8 CRC64;
Ouery Match
Best Local Similarity 100.0%; Pred. No. 6.7e+05;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps
Qy 4 LW 5
Db 8 LW 9
Search completed: February 12, 2003, 11:14:58
Job time: 10.7917 secs
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February 12, 2003, 11:10:52 ; Search time 12.5 Seconds
(without alignments)
53.300 Million cell updates/sec
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             908470 segs, 133250620 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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23
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                                                                                                                                                                                                                                                                                                                                            1 FXXLW 5
                                                                                                                                                                                                                                                                                     Title:
Perfect score:
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/SIDS2/gcgdata/geneseq/geneseqp-embl/AA1999.DAT:\*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2000.DAT:\*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2001.DAT:\*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2002.DAT:\* | SIDS2/gcddta/geneseq/geneseqp\_emb1/AA1989.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1990.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1992.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1992.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1992.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1994.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1994.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1996.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1996.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1996.DA1 | SIDS2/gcgdata/geneseq/geneseqp\_emb1/AA1996.DA1 

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		•			SUMMARIES	
Result		Que y				
No.	Score		Match Length DB	DB	ΙD	Description
	21	91.3	15	21	AAB29164	Peptide #8. Unide
7	20	87.0	S	20	AAY33038	Carbohydrate antiq
c	20	87.0	2	21	AAB07280	Motif binding to a
4	20	87.0	9	17	AAR89913	p53/MDM2 binding i
2	20	87.0	9	17	AAR89915	p53 minimal MDM2 b
9	20	87.0	9	18	AAW13606	p53 protein amino
7	20	87.0	9	21	AAB17074	Mdm/hdm antagonist
60	20	87.0	9	23	ABB73169	Mdm/hdm antagonist
6	20	87.0	7	17	AAR89921	Antibody DO-1 epit
10	20	87.0	7	17	AAR89925	p53 binding site k

Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide

FG loop sequence o TRAM-interaction m Ubiquitin binding Hepatitis GB virus Glycine max homogl Human oncogenic pr Amino acid sequenc Amino acid sequenc Amino acid sequenc Human p53 MDM2-con Mouse p53 MDM2-con p53 protein amino Human Leukocyte An Human Leukocyte An Human Leukocyte An Human Leukocyte An Human Leukocyte An Human Deukocyte An Human nocogenic pr Human oncogenic pr Human oncogenic pr Human oncogenic pr Mdm/hdm antagonist	0 00200
0 AAW94141 1 AAY57789 3 ABB09130 2 AAB62220 2 AAB62220 9 AAW46349 9 AAW46351 9 AAW46351 1 AAR89917 7 AAR89918 8 AAW13605 9 AAW37188 2 AAW37181 2 AAW37189 9 AAW37189 9 AAW37189 9 AAW37189	440mmmm40000
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## ALIGNMENTS

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Fork head associated, FHA; domain, transcriptional control, DNA replication; DNA repair; cell cycle control.
             AAB29164 standard; Peptide; 15 AA
                                                                                                                                         17-MAR-2000; 2000WO-GB01024.
                                                                                                                                                        99GB-0006432.
                                            (first entry)
                                                                                                                                                                                             Durocher D;
                                                                                                                                                                              (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                           WPI; 2000-664872/64.
                                                                                                            WO200057184-A2
                                                                                                                                                        19-MAR-1999;
28-JUN-1999;
                                                                                              Unidentified
                                                                                                                                                                                             Jackson SP,
                                                                                                                           28-SEP-2000.
                                           02-FEB-2001
                                                           Peptide #8
                             AAB29164;
RESULT 1
       AAB29164
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RESULT 3
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                                                                                                                                       molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA replication, and repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                                           The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these
                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mimotope; antigen; carbohydrate; antibody; mimic; vaccine; adjuvant; tumour immunity; cancer therapy; antibacterial; immune response; immunogenicity; anti-idiotype; T cell response manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Peptides and recombinant antibody mimics of carbohydrate antigens, used for, e.g. treatment of cancer and infection
                                                                                                                                                                                                                                                                                                                                                       91.3%; Score 21; DB 21; Length 15; 60.0%; Pred. No. 5.3e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Carbohydrate antigen peptide mimotope 11.
                                     Disclosure, Fig 2, 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 21; Page 72; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY33038 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US02405.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-NOV-1999 (first entry)
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                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 60.09
Matches 3; Conservative
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                                                                                                                                                                                                                                                                                                                 15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 FSALW 13
                                                                                                                                                                                                                                                                                                                                                                                                                                         1 FXXLW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-FEB-1998;
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                                                                                                                                                                                                                                                                                                                   Sequence
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tumour, meracrasis tequires ucracimment of maintains tumour, penetration of blood or lymph vessels and attachment to the endothelium of distant organs, ultimately resulting in the formation of new tumours. The selectin family of adhesion molecules is implicated in this process. E-selectin is a calcium-dependent molecule expressed by activated vascular endothelium. E-selectins bind to glycoconjugates or carrying a terminal tetrasaccharide Lewis antigen, which are found on tumour cell surfaces. One such Lewis antigen, which are found on tumour cell surfaces. One such Lewis antigen is Lewis Y (LeX). The binding of selectin molecules to their ligands is thought to be an important step in metastasis. Therefore, inhibition of E-selectin-dependent carbohydrate-mediated interactions is thought to be a target for anti-cancer therapy. The present sequence is a non planar-X-planar type motif. This motif is thought to bind to anti-LeY antibody BRSS-2. BRSS-2 is a peptido-mimetics of E-selectins. Peptides that block E-selectin-LeY binding and therefore adhesion of tumour cells and leukocytes to endothelial cells inhibit metastasis.
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                                                                                                                                                          amount of antigen required for immunization; (3) peptide synthesis may be more practical than synthesis of carbohydrate-protein conjugates or the production of anti-idiotypes; (4) peptide mimicking sequences can be engineered into DNA plasmids for DNA vaccination to further manipulate T cell responses. AAY33028-Y33087 represent the carbohydrate antigen peptide mimotopes described in the invention.
of the peptide mimics are; (1) the chemical composition and purity of synthesized peptides can be precisely defined; (2) the immunogenicity of the peptides can be significantly enhanced by polymerization or addition or relatively small carrier molecules that reduce the total
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptidomimetics of carbohydrate Lewis ligands useful for modulating inflammation, metastasis and angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            87.0%; Score 20; DB 20; Length 5; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; peptido-mimetic; tumour metastasis; E-selectin;
adhesion molecule; Lewis antigen; anti-adhesion therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Motif binding to anti-Lewis antigen anibody BR55-2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 37; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB07280 standard; peptide; 5 AA.
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Best Local Similarity 60.v.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200027420-A1.
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19-APR-1995;
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                     10-SEP-1996
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                                                                                                                                  01-FEB-1996.
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                                                                                         Synthetic.
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 AAR89915;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                       Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences ARR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alamine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                             New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
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                            87.0%; Score 20; DB 21; Length 5; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
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                                                                                                                                         AAR89913 standard; peptide; 6 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 29; 46pp; English.
                                                                                                                                                                                                    p53/MDM2 binding inhibitor #3
                                                                                                                                                                                                                                                                                                                  95WO-GB01719
                                                                                                                                                                                                                                                                                                                                     95US-0424957.
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                                                3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                       Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-105905/11.
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                                        Local Similarity
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          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 AA;
                                                                   1 FXXLW 5
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20-JUL-1994;
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                                                                                                                                                                                                                                                      Synthetic.
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Matches
                                                Matches
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AAR89915
ID AAR8
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                                                                                                                     RESULT 4
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AAR89915 standard; peptide; 6 AA.

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This is the sequence of the p53 tumour suppressor protein minimal MDM2 oncogene protein binding site. The sequence is used to generate peptides (ARA89911-3) conteg. this sequence, which forms part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos, and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
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                                                                                                      Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cpds, which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 15; 46pp; English.
                                                   p53 minimal MDM2 binding site #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW13606 standard; peptide; 6 AA.
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                                                                                                                                                                                                                                                                                                                                                            95WO-GB01719.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1996-105905/11.
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RESULT 8
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                                                                                                                                                                                                                                       The peptides AAW13602-6 represent peptide fragments derived from the wild type human p53 protein. This peptide corresponds to amino acids 18-23 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double minute-2 (mdm2) protein (AAW13600). Mdm2 protein is a 90 kD composed protein phich binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein cuppressor protein properties, especially in a p53-null background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoms of the colon, cancer of the breast, lung or stomach; myeloid leukaemia; B cell lymphoma, or other hyperproliferative conditions such as restenosis. Note: this sequence is not given in the specification but is constructed from the wild type human p53 sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified peptide, therapeutic agent, fusion, Fc domain, cancer, autoimmune disease, cytostatic, antiasthmatic, thrombolytic, VEGF, immunosuppressive, EPO, TPO, CTLA4, mimetic, IL-1, TNF, antagonist, MMP, inhibitor, erythropoietin, thrombopoietin, interleukin 1, cytotoxic T cell lymphocyte antigen 4, tumour necrosis factor, vascular endothelial growth factor, matrix metalloproteinase,
                                                                                                                                                                 Treating cancer with antagonist of oncogenic activity of protein Mdm2 - or nucleic acid encoding an antagonist, also viral vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.0%; Score 20; DB 18; Length 6; 60.0%; Pred. No. 7.88+05; 1ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mdm/hdm antagonist peptide sequence SEQ ID NO:130.
                                                                                                                 Dubs-Poterszman M, Tocque B, Wasylyk B;
                                                                           (INRM ) INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 asthma; thrombosis; pharmaceutical.
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                                                                                    (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                    Claim 4; Page -; 43pp; French.
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                         96WO-FR01340.
                                                  95FR-0010331.
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Best Local Similarity 60.0%
Matches 3; Conservative
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                                                                                                                                                                                          contg. this nucleic acid
                                                                                                                                        WPI; 1997-192837/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 AA;
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                         02-SEP-1996;
                                                  04-SEP-1995;
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 13-MAR-1997.
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The present invention describes composition of matter (1) comprising an FC domain, pharmacologically active peptides, and linkers. Where (I) is: (X1)a-F1-(X2)b, where: F1 = an FC domain; X1 and X2 = are each independently selected from -(IJ)c-F1-(IZ)d-F2.

-(IJ)c-P1-(IZ)d-P2-(IZ)e-P^3, or -(IJ)c-P1-(IZ)d-P2-(IZ)e-P3-(IA)F-P4 where P1, P2, P3, and P4 = are each independently sequences of pharmacologically active peptides; IJ, IZ, I3, and I4 = are each independently concert, provided that at least 1 of a and b is 1. The composition can have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can be used for producing pharmaceutical compositions are useful for treating cancer, asthma, thrombosis, or autoimmune diseases. The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions such as Fc receptor binding, prortein control in and possibly placental transfer. AAA69543 to AAAB86955 to AAAB803 represent nucleotide and amino acid sequences used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                   Novel composition of matter comprising an Fc domain and pharmacologically active peptides, useful for treating cancer and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                                                                                                                                                                  Liu C, Cheetham J, Boone TC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 39; Page 240; 608pp; English.
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98US-0105371.
99US-0428082.
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                                                                                                                                                                                                                                                                                                       WPI; 2000-350702/30.
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                                                                                                                              (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6 AA;
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23-OCT-1998;
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                                          22-OCT-1999;
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                                                                                                                                                                                                                       Feige U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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95US-0424957.
                                                                                                                                                                                                                                                                                                                              malignancies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-JUL-1994;
19-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FSDLW 5
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            20-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-1996
                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR89925;
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR8992!
g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ð
                                                                                                                                                                                                                                              multimers. (I) can have antiinflammatory, antitumour, immunosuppressive, cytostatic, antirheumatic, antiarthritic, antitumour, immunosuppressive, cytostatic, antirheumatic, antiarthritic, antitumour, immunosuppressive, cytostatic, anorectic, antiinfertility, haemostatic, dermatological, antiphactic activities. (I) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest, as a part of diagnostic kit to detect the presence of their proteins of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, theumatoid arthritis, diabetic retinopathy, obesity, sleep disorders, infertility, and neurological degenerative diseases. (I), comprising compounds are useful for treating disorders characterised by low red blood cell levels such as anaemia. The TPO-minetic comprising compounds are useful for treating conditions that involve an existing megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency is syndrome. Absolved an existing represent amino acid and nucleic acid sequences used in the expensent involve and abilisty represent involve and abilisty ceremplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus; antibody.
                                                                                                                                                                                                                                       The present invention describes a vehicle-peptide molecule (I) or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                          Novel vehicle-peptide molecule or its multimers useful for treating
                                                                                                                                                                    inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87.0%; Score 20; DB 23; Length 6; 60.0%; Pred. No. 7.8e+05; trive 0; Mismatches 2; Indels
                                                                                                        Gudas JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antibody DO-1 epitope on human p53 protein.
                                                                                                        Boone TC,
                                                                                                                                                                                                              Claim 39; Page 53; 176pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR89921 standard; peptide; 7 AA.
                                                                                                        Feige U, Liu C, Cheetham JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95WO-GB01719.
                            02-MAY-2001; 2001WO-US14310.
                                                     03-MAY-2000; 2000US-0563286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3; Conservative
                                                                                                                                  WPI; 2002-130313/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                               (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9602642-A1
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 08-NGV-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). Peptides contg. these sequence were identified by immobilising a library of peptides amino acids long) covering the mouse or human p53 protein and adding an extract contg. expressed MDM2 protein. The interaction was identified by the antibody D0-1. This is the sequence of a human p53 epitope for the antibody. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Interference, binding, p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus; antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
                                                                                                                                                                                                                                                         New cpds, which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               p53 binding site key amino acid for binding mouse MDM2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            87.0%; Score 20; DB 17; 60.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 27; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 26; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR89925 standard; peptide; 7 AA.
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94US-0277660.
94US-0277660.
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hes 3; Conservative
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                                                                                                                                    Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-105905/11.
                                                                   (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                                    WPI; 1996-105905/11.
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                                                                                                                                                                                                                                                                                                                                        other malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 AA;
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Sequence

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RESULT 11
                                                      AAW94141
X00000000000XX
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psi tumour suppressor protein and the WDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding macif between amino acids 16-33 (AAR89914). Peptides contg. these sequence were identified by immobilising a library of peptides (15 amino acids long) covering the mouse or human p53 protein and adding an extract contg. expressed MDM2 protein. This sequence represent the key amino acids in the p53 binding site required for mouse MDM2 binding. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies. New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncodene protein contain 0; Gaps 87.0%; Score 20; DB 17; Length 7; 60.0%; Pred. No. 7.8e+05; ative 0; Mismatches 2; Indels 3; Conservative Sest Local Similarity 7 AA; 1 FXXLW 5 3 FSDLW 7 Sequence Query Match Matches

FG loop sequence of ubiquitin-binding monobody clone 213. AAW94141 standard; peptide; 7 AA. 14-APR-1999 (first entry) AAW94141; 

Fibronectin type III; Fn3; monobody; beta-strand domain; loop region; specific binding partner; SBP; catalysis; LRS; ubiquitin.

Saccharomyces cerevisiae.

17-DEC-1998.

98WO-US12099 12-JUN-1998; 97US-0049410.

(RESE ) RESEARCH CORP TECHNOLOGIES INC.

Koide S;

WPI; 1999-060331/05.

Production of antibody compounds, particularly catalytic antibodies - using a fibronectin type III molecular scaffolding comprising beta-strand domain sequences and modified in one or more loop sednences

Example 10; Page 41; 96pp; English.

The invention relates to a synthetic fibronectin type III (Fn3) bolypeptide monobody that comprises Fn3 beta-strand domain sequences that are linked to loop region sequences (LRSs). One or more of the loop sequences in the synthetic Fn3 vary by deletion, insertion, or replacement of at least 2 amino acids from the corresponding LRss in wild-type Fn3. Host cells containing an expression vector comprising the synthetic Fn3 nucleic acid are used for the production of the Fn3 monobody. The invention also provides methods of identifying the amino acid sequence of a polypeptide molecule (i) capable of binding to a specific binding partner (SBP) so as to form a polypeptide:SBP complex, (ii) capable of catalysing a chemical reaction with a catalysed rate constant, Kcat, and an uncatalysed rate constant, Kucat, such that the ratio of the Kcat/Kunat is greater than 10. Sequences AAW44139-46 represent FG loop sequences of yeast ubiquitin-binding monobodies.

Score 20; DB 21; Length 7; Pred. No. 7.8e+05;

87.0%;

Query Match Best Local Similarity

Sequence

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                                                                                                                                                                                                                                                                                                                                                                      Transcriptional adapter motif; TRAM; TRAM interaction motif; TRIM; creb binding procein; transcriptional regulation; cytostatic; antiviral; cell cycle inhibitor; viral transcription inhibitor; cancer; tumour; viral disease; viral infection; cell cycle; apoptosis; growth arrest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for preparing a pharmaceutical composition and for disrupting an interaction between TRAM sequence and TRIM sequence in vitro, thereby inhibiting viral transcription or call cycle progression in mammalian cells especially cancer cell. Compounds which disrupt interaction between TRIM TRAM containing polypeptides can be used therapeutically to prevent or treat viral diseases and tumours. The polypeptides reduce susceptibility of cells to viral infection and regulate cell cycle antibodies against the TRIM or TRAM sequences. HPV types associated with high risk or low risk of cervical cancer can be distinguished with high risk or low risk of cervical cancer can be distinguished (CBP) TRAM sequence. The present sequence represents a TRIM sequence given in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a polypeptide comprising a transcriptional adaptor motif (TRAM) or a TRAM-interaction motif (TRAM) or a TRAM-interaction motif (TRAM) (I). (an be used for identifying compounds (II) comprising a polypeptide capable of disrupting an interaction between a TRAM sequence and/or a TRAM sequence or a TRAM sequence are identified by incubating, the polypeptide with (I) and determining if the polypeptide interacts with (I). Compound (II) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel polypeptides cells useful for treating viral disease and cancer
                                               Gaps
                                             .
87.0%; Score 20; DB 20; Length 7; 60.0%; Pred. No. 7.88+05; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                     TRAM-interaction motif derived from p53 SEQ ID NO:12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MOLE-) INST MOLECULAR & CELL BIOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 7; 73pp; English.
                                                                                                                                                                                                                 AAY57789 standard; peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99WO-GB01668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98GB-0011303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99GB-0000157,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Zimmermann H;
                                                                                                                                                                                                                                                                                           20-MAR-2000 (first entry)
                  Local Similarity 60.0 tes 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-072620/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O'Connor MJ,
                                                                                1 FXXLW 5
                                                                                                                    1 FADLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9961608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26-MAY-1999;
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                                                                                                                                                                                                                                                        AAY57789;
    Query Match
                                         Matches
                                                                                                                                                                            RESULT 12
                                                                                                                                                                                                AAY57789
                                                                                                                 Db
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Best Local Similarity
                         Hepatitis GB virus.
                                                                                                                                                                (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                            8 AA;
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                                                                                                                                                                                           Mushahwar IK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUN-2001
                                                                               07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Glycine max.
                                                                                                 14-FEB-1994;
13-MAY-1994;
                                                                                                                  29-JUL-1994;
23-NOV-1994;
                                             US6051374-A.
                                                              18-APR-2000
                                                                                                                                             30-JAN-1995
                                                                                                                                                                                                                                                          target HGBV
                                                                                                                                     23-NOV-1994
                                                                                                                                                                                  Dawson GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB62220;
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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AAB62220
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                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to fibronectin type III (Fn3) molecules comprising a stabilising mutation as compared to a wild-type Fn3. Fn3 can be used as a scaffold to engineer artificial binding proteins.

Modifications of the Fn3 scaffold that increase its stability are useful in that they allow the introduction of more mutations for better functions, and that these make it possible to use Fn3-based engineered proteins in a wider range of applications. The present sequence is a peptide described in the exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
Gaps
                                                                                                                                                                                                                                                                                                                                           New fibronectin type III molecule comprising a stabilizing mutation, useful for introducing more mutations for better functions, and in a wider range of applications
                                                                                                                                                         Fibronectin type 3; mutant; stabilising mutation; Fn3; antibody;
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Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis GB virus protein sequence SEQ ID NO:252.
5
Mismatches
                                                                                                                                                                                                                                                                             (RESE ) RESEARCH CORP TECHNOLOGIES INC. (KOID/) KOIDE S.
                                                                                                                                      Ubiquitin binding monobody 213 FG loop.
                                                                                                                                                                                                                                                                                                                                                                               Example 10; Page 138; 164pp; English.
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                                                                                 ABB94392 standard; Peptide; 7 AA.
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                                                                                                                    12-JUN-2002 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
 Conservative
                                                                                                                                                                                                                                                                                                                          WPI; 2002-171708/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match .
Best Local Similarity
Matches 3, Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7 AA;
                                                                                                                                                                  binding protein.
                                                                                                                                                                                                      WO200204523-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 FXXLW 5
                   1 FXXLW 5
                                    S
                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                       17-JAN-2002
3;
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                                    FSDLW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB09130;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                    ABB94392;
                                                                                                                                                                                                                                                                                                         Koide S;
Matches •′
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                                                               RESULT 13
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The present invention describe a method for detecting target hepatitis GB virus (HGBV) nucleic acid (THN) in a test sample (T) suspected of contraining HGBV. The method involves reacting (T) with a HGBV polynuclectide probe (I) containing 15 contiguous nuclectides, and which selectively hybridises to the HGBV genome or its full complement, and detecting the complex that contains THN, indicating the presence of target HGBV. The method is used for detecting target HGBV nucleic acid in the test sample suspected of containing HGBV and for characterisation of newly ascertained etiological agent of non-A, non-B, non-C, non-D and non-B hepatitis causing agents collectively termed as hepatitis GB virus. AAA55270 to AAA55489 and AAB089885 to AAB09480 represent nucleotide and protein sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Soybean, glutathione-S-transferase, GST, homoglutathione synthetase, HGS, beta-alanine, gamma-glutamylcysteine, herbicide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting target hepatitis GB virus nucleic acid in a test sample suspected of containing HGBV comprises reacting the test sample the HGBV polynucleotide probe and detecting the complex that contains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pilot-Matias TJ, Buijk S
Erker JC, Schlauder GG;
Hepatitis GB virus; HGBV; diagnosis; therapeutic; immunogenic;
infection; detection; characterisation; hepatitis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 9; Column 331-332; 369pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Muerhoff AS,
JN, Desai SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB62220 standard; peptide; 8 AA.
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94US-0242654.
94US-0283314.
94US-0344185.
94US-0344190.
95US-0377557.
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18-SEP-2000; 2000WO-GB03573.
           99GB-0022346.
               (ZENE ) ZENECA LTD.
          21-SEP-1999;
  29-MAR-2001.
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Townson JK, Edwards R, Cummins I, Skipsey M; Andrews CJ, Jepson I, WPI; 2001-257978/26.

sequences from soybean for producing plants which are resistant and tolerant to herbicide comprising fomesafen and/or acifluorfen Novel glutathione-S-transferase and homoglutathione synthetase

Claim 9; Page 39; 64pp; English.

The invention relates to new soybean glutathione-S-transferase (GST) and a homoglutathione synthetase (HGS). The HGS is capable of catalyzing the addition of beta-alanine onto gamma-glutamyloysteine. Polynucleotides encoding the proteins of the invention are useful for producing plants which are resistant and/or tolerant to a herbicide comprising fomesafen plants with further desired agronomic trait, especially resistant to a herbicide, comprising glyphosate or its salt. Further desired agronomic traits altered field, altered nutritional value, altered quality or any other desirable agronomic trait. GST or its variant is also useful as a selectable marker gene. The present sequence represents a Glycine max HGS protein fragment.

8 AA; Sequence

ö 0; Gaps Query Match

87.0%; Score 20; DB 22; Length 8; Best Local Similarity 60.0%; Pred. No. 7.8e+05; Matches 3; Conservative 0; Mismatches 2; Indels

1 FXXLW 5 δ

2 FAGLW 6 g

Search completed: February 12, 2003, 11:13:44 Job time : 13.5 secs

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Sequence 83, Appli
Sequence 4, Appli
Sequence 1, Appli
Sequence 36, Appli
Sequence 36, Appli
Sequence 252, App
Sequence 11, Appli
Sequence 11, Appli
Sequence 37, Appli
Sequence 38, Appli
Sequence 36, Appli
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                                                                                                        February 12, 2003, 11:15:02 ; Search time 3.3333 Seconds (without alignments) 38.323 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25,
Sequence 6, 7
Sequence 6, 7
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Sequence
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1: /cgn2_6/ptodata/1/pubpaa/USOB_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PUT_NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/PUT_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*
14: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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0 US-09-732-384-4

0 US-09-732-384-5

0 US-10-155-059-1

US-09-903-412-36

0 US-09-903-412-36

0 US-09-844-456-8

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-814-371-38

0 US-09-214-371-35

0 US-09-214-371-37

0 US-09-214-371-17

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                                                                                                                                                                                                                                                                                                                140259 segs, 25548876 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                     OM protein - protein search, using sw model
                                                                                                                                                                                                                                                         BLOSUM62
Gapop 10.0 , Gapext 0.5
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23
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Match Length DB
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887.00
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                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                            Run on:
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3, Appli
2, Appli
9, Appli
5, Appli
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                                                                                                                                                 Sequence 73, 7
Sequence 35636
Sequence 203, 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence:peptide, amin; OTHER INFORMATION: acid residues 18-23 of human p53
US-09-214-371-83
                                                                                      Sequence Sequence Sequence
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Sequence
Sequence
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0 US-09-214-371-39

0 US-09-214-371-40

0 US-09-214-371-40

0 US-09-214-371-74

0 US-09-214-371-74

0 US-09-214-371-74

0 US-09-732-384-7

2 US-10-155-059-13

2 US-10-155-059-10

2 US-10-155-059-10

2 US-10-155-059-15

2 US-10-155-059-16

2 US-09-214-371-73
                                                                                                                                                                          US-09-214-371-2
US-09-214-371-2
US-09-281-717-3
US-09-862-260A-9
US-09-881-276-5
US-09-881-276-6
US-10-155-059-21
US-10-155-059-21
                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garda-Echeverria, Carlos
APPLICANT: Borcal Echeverria, Carlos
APPLICANT: Garda-Echeverria, Carlos
APPLICANT: Garda-Echeverria, Carlos
APPLICANT: Garda-Echeverria, Carlos
APPLICANT: Garda-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interac
GURRENT APPLICATION NUMBER: U509/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                            ALIGNMENTS
                                                                                                                                                                                                                                                                                                                           ; Sequence 83, Application US/09214371B
; Patent No. US20010018511A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
   3; Conservative
    Query Match
Best Local Similarity
  US-09-214-371-83
   SEQ ID NO 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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; Sequence 4, Application US/09732384

RESULT 2 US-09-732-384-4

1 FXXLW | || 2 FSDLW

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                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Description of Artificial Sequence: Protein OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
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87.0%; Score 20; DB 10; Length 6; Best Local Similarity 60.0%; Pred. No. 1.2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h 87.0%; Score 20; DB 10; Length 6; Similarity 60.0%; Pred. No. 1.2e+05; 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Yuan, Zhi-Min
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, Jijia
TITLE OP INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 1999-12-07
PRIOR PILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
               GENERAL INFORMATION:
APPLICANT: Yuan, Zhi Min
APPLICANT: Yuan, Zhi Min
TILE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT APPLICATION NUMBER: 60/169,816
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTI VET. 2.1
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US-09-732-384-5
US-09-732-384-5
Sequence 5, Application US/09732384
Patent No. US20020132977A1
GENERAL INFORMATION
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                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
Patent No. US20020132977A1
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                             COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Koide, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY FOLYPEPTIDES
FILE REFERENCE: 109.050US1
CURRENT APPLICATION NUMBER: US/09/903,412
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: US 60/217,474
PRIOR PILING DATE: 2000-07-11
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     87.0%; Score 20; DB 9; I
60.0%; Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION - UNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
FILING DATE: 12-May-1997
FILING DATE: 12-May-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 36, Application US/09903412; Publication No. US20030027319A1; GENERAL INFORMATION:
                                                                                                                                                         ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: (Unknown) INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3; Conservative
                                                                                                             STATE: MA
COUNTRY: USA
                                                                                            CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Best Local Similarity
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US-09-903-412-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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APPLICANT: JAMES C. BERER
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
STATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  87.0%; Score 20; DB 8; Length 8; 60.0%; Pred. No. 1.2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                      COMPACT: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
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APPLICANT: Houghton, Michael
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
APPLICANT: Rio, George
TITLE OF INVENTION: Hoperitis C virus protease
FILE REFERENCE: 223002010005
CURRENT APPLICANTON NUMBER: US/09/884,456
CURRENT FILING DATE: 1990-06
PRIOR APPLICATION NUMBER: 08/253,230
PRIOR APPLICATION NUMBER: 08/219
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/440,548
PRIOR FILING DATE: 1995-05-12
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR APPLICATION NUMBER: 07/680,296
PRIOR APPLICATION NUMBER: 07/680,296
PRIOR APPLICATION NUMBER: 07/505,433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435435
ATTORNEY/ACENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECHONI: 708-937-6365
TELEPHONE: 708-938-2623
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NUMBER OF SEQ ID NOS: 90
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-884-456-8
; Sequence 8, Application US/09884456
; Publication No. US20030027317A1
; GENERAL INFORMATION:
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Best Local Similarity 60.0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) MOLECULE TYPE: protein US-08-424-5508-252
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                                                                     2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A. STREET: 121 South Eighth Street, Ste. 1600 CITY: Minneapolis
                                                                                                                                                                                                                                                                  Sequence 36 Application US/09096749A
Patent No. US2002019317A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Koieda, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Ann S. Vikenins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 252, Application US/08424550B
Patent No. US20020119447A1
GENERAL INFORMATION:
APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURBSH M. DESAI
APPLICANT: THOMAS P. LEARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/096,749A FILING DATE: June 12, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTEQ Version 2.0b
CURRENT APPLICATION DATA:
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                                                          Matches ' 3; Conservative
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STRANDEDNESS: single
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FRAGMENT TYPE: i
ORIGINAL SOURCE:
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Gaps . 0

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87.0%; Score 20; DB 9; Length 9;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
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87.0%; Score 20; DB 9; Length 9;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                      FEATURE:

COTHER INFORMATION: West Nile Fever virus protease
US-09-884-456-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Houghton, Michael
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
TITLE GOT INVENTION: Hoperitis C virus protease
FILE REFERENCE: 223002010005
CURRENT APPLICATION NUMBER: US/09/884,456
CURRENT FILING DATE: 1990-02-19
PRIOR FILING DATE: 1996-09-06
PRIOR APPLICATION NUMBER: 08/709,177
PRIOR FILING DATE: 1996-09-06
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR PRILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR FILING DATE: 1900-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; FEATURE:
; OTHER INFORMATION: Murray Valley virus protease
US-09-884-456-11
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APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, George
ITILE OF INVENTION: Hepatitis C virus protease
FILE REFERENCE: 223002010005
CURRENT APPLICATION NUMBER: US/09/884,456
CURRENT PILING DATE: 2001-06-18
PRIOR APPLICATION NUMBER: 09/253,230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 11, Application US/09884456; Publication No. US20030027317A1; GENERAL INFORMATION:
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                                                      TYPE: PRT ORGANISM: West Nile Fever virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Murray Valley virus
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US-09-884-456-11
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US-09-884-456-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Bottger, Aggelica
APPLICANT: Bottger, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TILLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

87.0%; Score 20; DB 10; Length 9;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 87.0%; Score 20; DB 9; Length 9; Best Local Similarity 60.0%; Pred. No. 1.2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
             PRIOR PELICATION NUMBER: 08/40,177
PRIOR FILING DATE: 1996-09-06
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR FILING DATE: 1995-05-12
PRIOR PLING DATE: 1995-05-12
PRIOR PLING DATE: 1995-05-12
PRIOR APPLICATION NUMBER: 08/350,884
PRIOR PILING DATE: 1994-12-06
PRIOR PLING DATE: 1991-04-04
PRIOR PLING DATE: 1991-04-04
PRIOR PLING DATE: 1990-04-04
PRIOR PLING DATE: 1990-04-04
PRIOR PLING DATE: 1990-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR FILING DATE: 1900-04-04
                                                                                                                                                                                                                                                                                                                                                                                                                                           ), OTHER INFORMATION: Kunjin virus protease
US-09-884-456-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 37, Application US/09214371B ; Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: X = Ac=Cys(Acrd)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lane, David
Bottger, Volker
Bottger, Angelica
PRIOR FILING DATE: 1999-02-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: VARIANT LOCATION: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: VARIANT LOCATION: (9)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 37
LENGTH: 9
                                                                                                                                                                                                                                                                                                                      SEQ ID NO 14
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                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Puret, Pascal III, Carlos APPLICANT: Puret, Pascal III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III, Carlos III,
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NAME/KEY: VARIANT
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OTHER INFORMATION: Description of Artificial Sequence:peptide
NAME/KEY: VARIANT
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                                                                                                                                                   Bottger, Volker
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Bcheverria, Carlos
                                  Sequence 38, Application US/09214371B
Patent No. US20010018511A1
GENERAL INFORMATION:
APPLICANT: Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 35, Application US/09214371B Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; LOCATION: (9)
; OTHER INFORMATION: X = Pro-NH2
US-09-214-371-38
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US-09-214-371-38
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...rulCANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Ehreveria, Carlos
APPLICANT: Firet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20937/A/POT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: PCT/EP97/03549
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR PILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
TYPE: pr
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                                                                                                                                        87.0%; Score 20; DB 10; Length 10; 60.0%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.0%; Score 20; DB 10; Length 10; 60.0%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
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; Patent No. USZ0010018511A1

; GENERAL INFORMATION:

; APPLICANT: Lane, David

; APPLICANT: Bottger, Volker

; APPLICANT: Bottger, Nagelica

; APPLICANT: Picksley, Stephen

; APPLICANT: Chene, Patrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : Lane, David

: Bottger, Volker

: Bottger, Angelica

: Picksley, Stephen

Chene, Patrick

Hochkeppel, Heinz-Kurt

Garcia-Echeverria, Carlos

Furet, Pascal
LOCATION: (1)
OTHER INFORMATION: X = Ac-Cys(Acrd)
NAME/KEY: VARIANT
                                                          LOCATION: (10)
COTHER INFORMATION: X = Pro-NH2
US-09-214-371-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: (10); OTHER INFORMATION: x = Pro-NH2 US-09-214-371-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: VARIANT
LOCATION: (1)
OTHER INFORMATION: X = AC-CYS
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ORGANISM: Artificial Sequence
                                                                                                                                        Query Match 87.0
Best Local Similarity 60.0
Matches 3; Conservative
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Best Local Similarity
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Sequence 16, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 114, App
Sequence 11, Appl
Sequence 11, Appl
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                                                                                                                                                         February 12, 2003, 11:13:52; Search time 99.375 Seconds (without alignments) 32.439 Willion cell updates/sec
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1: /cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*
2: /cgn2 6/ptodata/1/paa/USOG COMB.pep:*
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GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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US-09-936-956-10

PCT-US99-02405-11

PCT-US99-26277-114

US-09-601-558-11
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Maximum Match 100%
Listing first 45 summaries
                                                                                                        - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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23
1 FXXLW 5
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Match Length
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Maximum DB seq length: 25
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91.3
87.0
87.0
87.0
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                                                                                                           OM protein
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                                                                                                                                                              Run on:
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Sequence 114, App
Sequence 11, Appl
Sequence 2, Appli
Sequence 2, Appli
Sequence 83, Appli
Sequence 83, Appli
Sequence 130, App
                                                                                                                                               Sequence 5, Appli
Sequence 1, Appli
Sequence 27, Appl
Sequence 36, Appl
Sequence 36, Appl
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Sequence 604, App
Sequence 381, App
Sequence 123, App
Sequence 123, App
Sequence 123, App
Sequence 252, App
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Sequence 130, App
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                                                                                                                                    Sequence 4, Appl
                                                                                                                       Sequence 130,
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Sequence 304,
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Sequence
Sequence
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US-08-277-660-16
Sequence 16, Application US/08277660
GENERAL INFORMATION:
APPLICANT: Dicksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIREE: Four Emparated Center, State 3400
CITY: San Francisco
COUTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
CURRENT APPLICATION NUMBER: US/08/277,660
FILING DATE: 20-10L-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: DISGREY, WALTER H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
2 US-09-831-047C-114
2 US-09-84-594-11
US-08-277-660-2
US-08-277-660-3
6 US-09-214-130
US-08-277-660-3
6 US-09-214-130
9 US-09-214-130
9 US-09-563-286B-130
9 US-09-563-286B-130
9 US-09-563-286B-130
1 US-09-732-384-5
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8 US-08-446-755-8
8 US-08-486-749-252
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8 US-08-486-745-252
8 US-08-486-745-252
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US-09-458-297-345
US-09-458-297-475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Trustees of the University of Pennsylvania TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens FILE REPERENCE: upn3544 CURRENT APPLICATION NUMBER: PCT/US99/02405 CURRENT FILING DATE: 1999-02-04 EARLIER APPLICATION NUMBER: 60/073,690 EARLIER, FILING DATE: 1998-02-04 NUMBER OF SEQ ID NOS: 60 SOFTWARE; Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KuDOS Pharmaceuticals Limited
APPLICANT: KuDOS Pharmaceuticals Limited
APPLICANT: Jackson, Stephen P
APPLICANT: Jackson, Stephen P
TITLE OF INVENTION: Screening Methods Based on FHA Domains
FILE REFERENCE: MEWBOLK
CURRENT APPLICATION NUMBER: US/09/936,956
CURRENT APPLICATION NUMBER: GB 9906432.1
PRIOR FILING DATE: 1999-00-19
PRIOR FILING DATE: 1999-06-28
PRIOR FILING DATE: 1999-06-28
NUMBER OF SEQ ID NOS: 57
SOFFWARE: PATENTIN VET: 2.1
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                                                                                                                                                                                                                                                                                                     Query Match 91.3%; Score 21; DB 6; Length 11; Best Local Similarity 60.0%; Pred. No. 6.1e+03; Matches 3; Conservative 0; Mismatches 2; Indels
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                TELEPHONE: (415) 781-1969
TELERY: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acids
TELECOMMUNICATION INFORMATION:
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                                              ; OTHER INFORMATION: Description of Artificial Sequence: Novel Sequence PCT-US99-02405-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Maddalena
APPLICANT: Keiber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of CancerFILE REPERSNCE: WST93PCT
CURRENT APPLICATION NUMBER: PCT/US99/26277
CURRENT APPLICATION NUMBER: 60/107,478
EARLIER FILING DATE: 1999-11-06
EARLIER FILING DATE: 1999-11-06
WIUMBER OF SEQ ID NOS: 12.0
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APPLICANT: The Trustees of the University of Pennsylvania
TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens
TITLE OF INVENTION: PEPTIGE WIMOTOPES OF CARBOHY FILE REFERENCE: UPN3567
CURRENT APPLICATION NUMBER: US/09/601,558
CURRENT APPLICATION NUMBER: PCT/US99/02405
PRIOR PLING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 60
SEQ ID NOS: 80
SEQ ID NO 11
LENGTH: 5
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 114, Application PC/TUS9926277; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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LENGTH: 5
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                        FEATURE:
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i Sequence 114, Application US/09831047C

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sequence 114, Application US/09831047C

septicant: The Wistar Institute of Anatomy and Biology

APPLICANT: The Wistar Institute of Anatomy and Biology

APPLICANT: The Mistar Institute of Anatomy and Biology

TITLE OF INVENTY: Kieber-Emmons, Thomas

TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer;

FILE REPRENCE: WST93AUSA

CURRENT APPLICATION NUMBER: US/09/831,047C

CURRENT APPLICATION NUMBER: 60/107,478

PRIOR APPLICATION NUMBER: 60/107,478

PRIOR FILING DATE: 1998-11-06

NUMBER OF SEQ ID NOS: 121

SEQ ID NO 114

LENGTH: 5
                                                                                                                                                                                                                                          Sequence 114, Application US/09831047B
; Sequence 114, Application US/09831047B
; GENERAL INFORMATION:
; APPLICANT: The Wister Institute of Anatomy and Biology
; APPLICANT: The Trustees of the University of Pennsylvania
; APPLICANT: Baszczyk-Thurin, Magdalena
; APPLICANT: Homes
; TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer
; FILE REPRENCE: WST93AUSA
; CURRENT FILING DATE: 2001-08-14
; PRIOR APPLICATION NUMBER: 60/107,478
; PRIOR APPLICATION NUMBER: 60/107,478
; RILING DATE: 1998-11-06
; NUMBER OF SEQ ID NOS: 121
; SOFTWARE: PatentIn Ver. 2.0
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Score 20; DB 20; Length 5; Pred. No. 4.2e+06; 0; Mismatches 2; Indels
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CTHER INFORMATION: peptido-mimetic of a Lewis antigen
US-09-831-047B-114
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  87.0%;
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                                                      Conservative
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Best Local Similarity
     Query Match
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Matches 3; Conserv
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APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens and DNA Molecules Encorntries of INVENTION: Same
FILE REFERENCE: UDN-3984
CURRENT APPLICATION NUMBER: US/09/894,594
CURRENT APPLICATION NUMBER: 09/601,558
PRIOR APPLICATION NUMBER: 09/601,558
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
PRIOR FILING DATE: 1998-02-04
PRIOR FILING DATE: 1998-02-04
PRIOR FILING DATE: 1900-06-28
NUMBER: OF SEQ ID NOS: 70
SEQ ID NO 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: FEIGE, ULRICH
APPLICANT: GHEETHAM, JANET
APPLICANT: BOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20; DB 22; Length 5;
Pred. No. 4.2e+06;
0; Mismatches 2; Indels
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CURRENT APPLICATION NUMBER: PCT/US99/25044
CURRENT FILING DATE: 1999-10-25
EARLIER PILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE: Patentin Ver. 2.1
SEG ID NO 130
LENGTH: 6
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RESULT 8
US-09-894-594-11
Sequence 11, Application US/09894594
; GENERAL INFORMATION:
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; OTHER INFORMATĪON: Novel Sequence
US-09-894-594-11
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60.0%;
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Matches 3; Conservative
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                                         Sequence 2, Application US/08277660
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Picksley, Steven M.
TITLE OF INVENTION: Unterruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 3, Application US/08277660
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
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87.0%; Score 20; DB 6; Length 6;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STRET: Four Embarcadero Center, Suite 3400
CITY: Brancisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660 FILING DATE: 20-JUL-1994 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                              ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                          COUNTRY: United States
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 6 amino acids TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-277-660-2
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                      US-08-277-660-2
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US-08-277-660-3
RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lane, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeveria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20937/APCT
CURRENT PILLING DATE: 1999-03-26
FRICH APPLICATION NUMBER: US/09/214,371B
CURRENT FILLING DATE: PCT/EP97/03549
FRICH PILLING DATE: 1997-07-04
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US-09-214-371-83
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87.0%; Score 20; DB 16; Length 6;

Best Local Similarity 60.0%; Pred. No. 4.2e+06;

Matches 3; Conservative 0; Mismatches 2; Indels
                                          FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: A-60244/WHD
TELECOWUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELERAX: (415) 398-3249
TELERAX: (415) 781-1989
TELERAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                   APPLICATION NUMBER: US/08/277,660
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-214-371-83; Sequence 83, Application US/09214371B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 130, Application US/09428082;
GENERAL INFORMATION:
APPLICANT: LIU, CHUAN-FA
APPLICANT: FEIGE, ULRICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 83
LENGTH: 6
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                 LENGTH: 6 amino acids
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; Gequence 130, Application US/09428082B
; GENERAL INFORMATION:
; APPLICANT: ELIG. CHUAN-FA
; APPLICANT: CHEETHAM, JANET C.
; APPLICANT: BOONE, THOMAS CHARLES
; TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
; TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
; TITLE OF INVENTION WHERE: US/09/428,082B
; CURRENT FILING DATE: 1999-10-22
; PRIOR APPLICATION NUMBER: 60/105,371
; PRIOR FILING DATE: 1999-10-23
; NUMBER OF SEQ ID NOS: 1133
; SOFTWARE: Patentin version 3.1
; FENCEN: APPLICATION VERSION 3.1
; APPLICANT: CHEETHAM, JANET
; APPLICANT: BOONE, THOMAS CHARLES
; TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
; FILE REFERENCE: A-5.7
; CURRENT PILICATION NUMBER: US/09/428,082
; CURRENT PILICATION NUMBER: G0/105,371
; EARLIER APPLICATION NUMBER: 60/105,371
; EARLIER PILICATION NUMBER: 60/105,371
; EARLIER PILICATION NUMBER: 1998-10-23
; NUMBER OF SEQ ID NOS: 1133
; SOFTWARE: PALENTIN Ver. 2.1
; SEQ ID NO 130
; LENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                ) OTHER INFORMATION: Description of Artificial Sequence:MDM/HDM; OTHER INFORMATION: ANTAGONIST PEPTIDE US-09-428-082-130
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Sequence 130, Application US/09563286B
Sequence 130, Application US/09563286B
GENERAL INFORMATION:
APPLICANT: FEIGE, ULRICH
APPLICANT: CHESTHAM, JANET C.
APPLICANT: GENERAL JANET C.
APPLICANT: GUDS, JEAN MARIE
TITLE OF INVENTION: WODIFIED PEPTIDES AS THERAPEUTIC AGENTS
FILE REFERENCE: A-527A
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Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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87.0%; Score 20; DB 18; Length 6;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
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87.0%; Score 20; DB 19;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Mdm/hdm antagonist peptide US-09-563-286B-130
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                PRIOR APPLICATION NUMBER: 09/428,082
PRIOR FILING DATE: 1999-10-22
PRIOR APPLICATION NUMBER: 60/105,371
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1151
SOFTWARE: PatentIn version 3.1
CURRENT FILING DATE: 2000-05-03
                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            1 FXXLW 5
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                                                                                                                                                     SEQ ID NO 130
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Sequence 36, Appl
Sequence 286, App
Sequence 27, Appl
Sequence 27, Appl
Sequence 141, Appl
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9, Appli
343, App
12722, A
3, Appli
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2196, Ap
1920, Ap
1994, App
12994, A
115, App
149, App
149, App
155, App
155, App
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Sequence 36, Appl
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                                                                                               (without alignments)
59.144 Million cell updates/sec
                                                                               February 12, 2003, 11:14:12; Search time 7.08333 Seconds
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1: /cgn2 6/ptodata/1/paa/USOG_NEW COMB.pep:*
2: /cgn2 6/ptodata/1/paa/USOG_NEW COMB.pep:*
3: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
7: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
7: /cgn2 6/ptodata/1/paa/USOB_NEW COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-10-174-717A-36
US-09-701-080C-22
US-09-701-080C-27
US-06-438-805-9
US-10-211-088-141
US-60-438-805-9
US-10-203-138A-1272
US-10-225-567A-2041
US-10-225-567A-2041
US-10-225-567A-199
US-10-225-567A-199
US-10-225-567A-199
US-10-235-567A-2196
US-10-235-567A-199
US-10-235-567A-199
US-10-235-567A-199
US-10-235-567A-199
US-10-235-567A-199
US-10-235-567A-199
US-10-235-59-139
US-10-283-599-139
US-10-283-599-139
US-10-283-599-139
US-10-283-599-139
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US-10-283-599-149
US-10-283-599-149
US-10-283-144-37
US-10-283-144-37
US-10-283-144-37
US-10-283-144-37
US-10-283-144-37
US-10-171-734-4
US-10-099-056-2094
                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                    OM protein - protein search, using sw model
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23
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Match Length
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Maximum DB seq length: 25
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US-09-701-080C-12

Sequence 12, Application US/09701080C

Sequence 12, Application US/09701080C

Sequence 12, Application US/09701080C

GENERAL INFORMATION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 17 TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION

FILLE REFERENCE: N3447C GCW

CURRENT APPLICATION NUMBER: US/09/701,080C

CURRENT FILLING DATE: 1998-06-26

PRIOR FILLING DATE: 1998-06-26

PRIOR FILLING DATE: 1998-01-05

PRIOR FILLING DATE: 1999-01-05

NUMBER OF SEQ ID NOS: 36

NUMBER OF SEQ ID NOS: 36
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1318, Ap
316, App
316, App
13, App
12, App
12, App
312, App
372, App
380, App
372, App
372, App
372, App
16, App
173, App
16, App
173, App
173, App
173, App
180, App
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NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, St. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87.0%; Score 20; DB 5; Length 7; 60.0%; Pred. No. 3.5e+05; Live 0; Mismatches 2; Indels
                 PCT-USG2-34021-316
PCT-USG2-34021-399
US-10-282-958-20
US-10-282-958-20
US-10-282-958-39
PCT-USG2-24317-31
US-00-62-109A-372
US-10-062-109A-372
US-10-062-109A-372
US-10-062-109A-372
US-10-062-109A-372
US-10-062-109A-372
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US-10-062-109A-73
US-10-062-109A-73
US-10-062-109A-73
US-10-062-109A-73
US-10-062-109A-73
US-10-062-109A-73
US-10-062-109A-73
US-10-269-171A-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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ZIP: 55402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 3; Conserv
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  1 FXXLW 5
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US-10-174-717A-36
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1 FXXLW 5
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US-09-701-080C-27
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LENGTH: 13
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LENGTH: 13
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APPLICANT: Staney, John
APPLICANT: Staney, John
APPLICANT: Staney, John
APPLICANT: Staney, W. Martin
APPLICANT: Epimmune, Inc.
TITLE OF INVENTION: HLA Binding Peptides and Their Uses
TILE REFERENCE: 39963-20019.20
CURRENT FILING DATE: 1998-11-10
PRIOR PILING DATE: 1998-11-10
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SEQ ID NO 286
LENGTH: 11
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87.0%; Score 20; DB 5; Length 11;
Best Local Similarity 60.0%; Pred. No. 4e+02;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           87.0%; Score 20; DB 6; Length 7; 60.0%; Pred. No. 3.5e+05; tive 0; Mismatches 2; Indels
             SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/174,717A
FILING DATE: 18-Jun-2002
PRIOR APPLICATION DATA:
RAPLICATION DATA:
APPLICATION NUMBER: 09/096,749
FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY/ABONT INFORMATION:
NAME: Ann S. VİKSHİNS
REGISTRATION NUMBER: 37,748
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO FRAGMENT TYPE: internal ORIGINAL SOURCE: SEQUENCE DESCRIPTION: SEQ ID NO: 36: US-10-174-717A-36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 286, Application US/09189702A GENERAL INFORMATION:
OPERATING SYSTEM: WINDOWS
                                                                                                                                                                                                                                                                                                                               TELEFAX: (612) 339-3061
                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
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Matches 3; Conservative
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KENDULA WE SECULATION OF SECULATION OF SECULAR AND CELL BIOLOGY CENERAL INPORMATION:

SEQUENCE 22 APPLICATION OF MOLECULAR AND CELL BIOLOGY CENERAL INPORMATION:

TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTISIN P300 INTITLE OF INVENTION: PRANCRIPTIONAL REGULATION

TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTISIN P300 INTITLE OF INVENTION: PRANCRIPTIONAL REGULATION

CURRENT APPLICATION NUMBER: GW 9911303.8

PRIOR FILING DATE: 1998-05-26

PRIOR FILING DATE: 1999-0157.0

PRIOR FILING DATE: 1999-0157.0

NUMBER OF SEQ ID NOS: 36

SOFTWARE: PATENTIN VET. 2.1
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GENERAL INFORMATION:
TATLE OF INVENTION: POLYPERTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 ITILE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILLE REFERENCE: N7347C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: GB 9811303.8
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR PLILING DATE: 1998-05-26
PRIOR PLILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87.0%; Score 20; DB 5; Length 13; 60.0%; Pred. No. 4.5e+02; Lindels o; Mismatches 2; Indels
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APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
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Matches 3; Conservative
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GENERAL INFORMATION:
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Molecular David K.
APPLICANT: Rank, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: HANALYSIS OF GENE EXPRESSION IN BT 474
FILES OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
FILE REFERENCE: PB 0004 WO 8
CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PLILNG DATE: 26 May 2000 (04.02.00)
PRIOR FILING DATE: 03 August 2000 (03.08.00)
PRIOR FILING DATE: 03 August 2000 (03.08.00)
PRIOR PLILNG DATE: 03 Cotober 2000 (03.10.00)
PRIOR PLILNG DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 21 September 2000 (21.09.00)
PRIOR FILING DATE: 30 June 2000 (30.06.00)
NUMBER OF SEQ ID NOS: 15438
SOFTWARE: Molecular Dynamics Sequence Listing Engine
SEQ ID NO 12722
LEMOTH: 22
LEMOTH: 22
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US-10-203-138A-12722
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

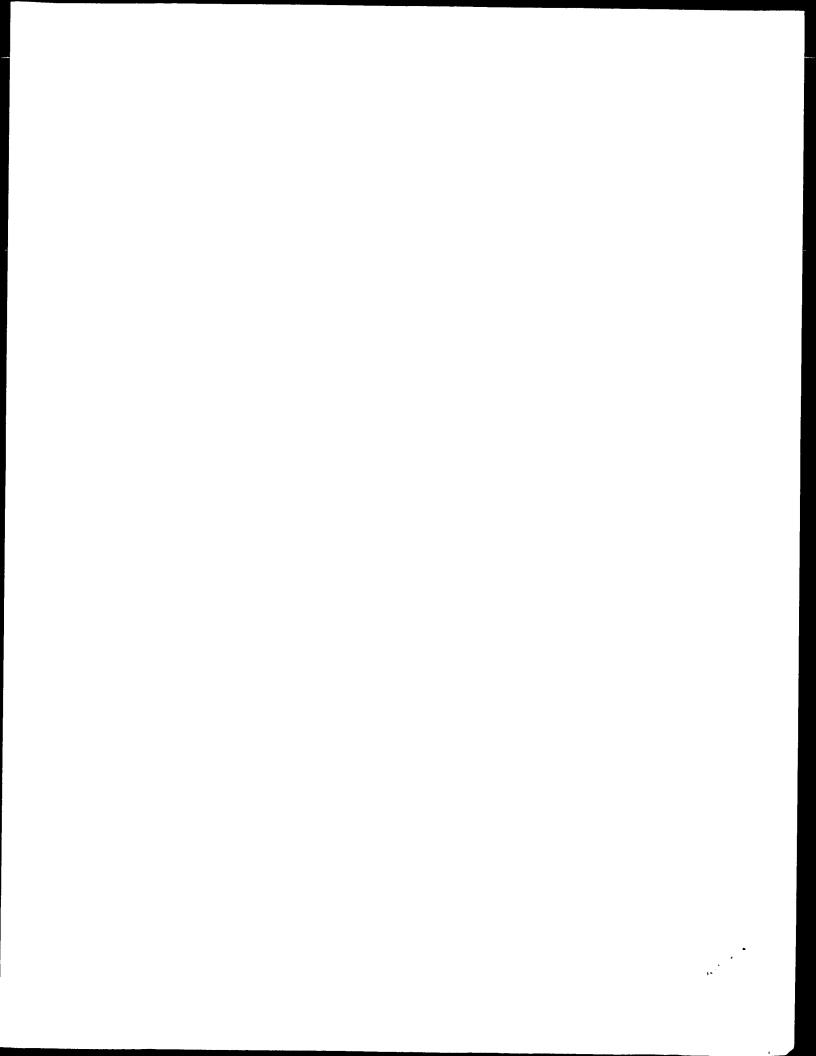
87.0%; Score 20; DB 6; Length 22;
Best Local Similarity 60.0%; Pred. No. 6.9e+02;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                 87.0%; Score 20; DB 1; Length 20; 60.0%; Pred. No. 6.4e+02;
                                                                                                                                                                                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL =
                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12722, Application US/10203138A GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT: Baxter, John
APPLICANT: Darimont, Beatrice
; APPLICANT: Feng, Weijun
                                        NAME/KEY: MISC FEATURE
) OTHER INFORMATION: DGI7-20M-4-G12
PCT-US02-34021-343
                                                                                                                                                                                                    3; Conservative
ORGANISM: Eschericia coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
                                                                                                                                                                         Best Local Similarity
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US-10-203-138A-12722
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                                                                                                                                                      Query Match
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: TARGET SPECIFIC SCREENING AND ITS USE FOR IDENTIFYING TARGET BIND:
FILE REFERENCE: 2598-4009PC
CURRENT APPLICATION NUMBER: PCT/US02/34021
CURRENT APPLICATION NUMBER: PCT/US02/34021
PRIOR FILING DATE: 2002-10-24
PRIOR FILING DATE: 2001-10-24
NUMBER OF SEQ ID NOS: 537
SOFTWARE: PALENTIN Version 3.1
SEQ ID NO 343
LENGTH: 20
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                     TITLE OF INVENTION: NOVEL Fusion Proteins And Assays For Molecular Binding FILE REFERENCE: 01-1022-US
CURRENT APPLICATION NUMBER: US/10/211,088
CURRENT ETLING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/309,395
PRIOR FILING DATE: 2001-08-01
PRIOR FILING DATE: 2001-08-01
PRIOR FILING DATE: 2001-12-13
NUMBER OF SEQ ID NOS: 366
SOFTWARE: 2001-12-13
LENGTHARE: PATENTIN VERSION 3.1
LENGTHARE: PATENTIN VERSION 3.1
LENGTH: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; ; OTHER INFORMATION: Peptide
US-60-438-805-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 9, Application US/60438805; GENERAL INFORMATION: THOMAS; APPLICANT: KOADAEK, THOMAS; TITLE OF INVENTION: MIXED ELEMENT CAPTURE AGENTS; FILE REFERENCE: UTSD:935USP1.
CURRENT APPLICATION NUMBER: US/60/438,805; CURRENT FILING DATE: 2003-01-09; NUMBER OF SEQ ID NOS: 12; SOFTWARE: Patentin Ver. 2.1; SEQ ID NO 9.
                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Binding domain US-10-211-088-141
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  Chen, Yih-Tai
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US-60-438-805-9
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GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sexul, Renate
APPLICANT: Sexul, Rabelle
APPLICANT: Cornut, Isabelle
APPLICANT: MICONICQ, Jean
TITLE OF INVENTION: GENE THERAPY APPROACHES TO
TITLE OF INVENTION: SUPPLY APOLIPOPROTEIN A-1 AGONISTS AND THEIR
TITLE OF INVENTION: USE TO TREAT DYSLIPIDEMIC DISORDERS.
NUMBER OF SEQUENCES: 274
                                                                                                                                                                                                                                                                                                                                          ..
                                                                                                                                                                                                                                                                                         82.6%; Score 19; DB 6; Length 14; 60.0%; Pred. No. 7.7e+02; Live 0; Mismatches 2; Indels
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                  CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
PRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 2196
LENGTH: 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 1920, Application US/10225567A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 1920
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 60.09
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         FILE REFERENCE: 1920-4-4
                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-225-567A-2196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Homo sapiens
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 13
US-10-225-567A-1920
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US-10-283-599-199
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APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Brumer, Glenna C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORG FILE OF INVENTION NUMBER: US/10/225,567A
CURRENT PELING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 60/257,144
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
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GENERAL INFORMATION:
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
                                                  APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR
TITLE OF INVENTION: COACTIVAGR BINDING
FILE REFERENCE: 9811-008-999
CURRENT APPLICATION NUMBER: US/09/281,717A
CURRENT AILING DATE: 1999-03.30
CRIENT APPLICATION NUMBER: US 60/079,956
PRIOR APPLICATION NUMBER: US 60/079,956
PRIOR PILING DATE: 1998-03.30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LOCATION: (2)...(3)
; OTHER INFORMATION: Xaa is any amino acid US-09-281-717A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2041, Application US/10225567A GENERAL INFORMATION:
  Fletterick, Robert
                       Kushner, Peter
West, Brian
Wagner, Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
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Matches 5; Conserv
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LENGTH: 12
APPLICANT:
                    APPLICANT:
APPLICANT:
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APPLICANT: LifeSpan Biosciences
APPLICANT: Enven, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
APPLICANT: Roush, Christine L.
APPLICANT: Roush, Christine L.
APPLICANT: ROUSH, Christine L.
APPLICANT ROUSH, Christine L.
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
PRIOR FILING DATE: 2001-12-19
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APPLICATION NUMBER: US 08/209,762
FILING DATE: 11-MAR-1995
ATTORNEY AGENT INFORMATION:
NAME: MCDORNELL, JOHN J.
REGISTRATION NUMBER: 26,949
REFERENCE/DOCKET NUMBER: 93,1133-J
TELECOMMUNICATION INFORMATION:
TELEPONE: 312-715-1000
TELEFAX: 312-715-1000
TELEFAX: 312-715-1234
TELEX: 910-221-5317
INFORMATION FOR SEQ ID NO: 54:
SEQUENCE CHARACTER:STICS:
                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc feature
OTHER INFORMATION: "BPI.7"
SEQUENCE DESCRIPTION: SEQ ID NO: 54:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Search completed: February 12, 2003, 11:24:44 Job time: 7.08333 secs
                                                                                                                                                                                                                                                                     LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 FLKLW 12
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US-10-319-786-54
; Sequence 54, Application US/10319786
; Sequence 54, Application US/10319786
; GENERAL INFORMATION:
; APPLICANT: Little,
; TITLE OF INVENTION: Biologically Active Peptides from
; TITLE OF INVENTION: Protein and Uses Thereof
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COMPUTER: 1BM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Rclease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/319,786
FILING DATE: 13-Dec-2002
PRIOR APPLICATION NUMBER: US/08/473,344
FILING DATE: 7-JUN-1995
APPLICATION NUMBER: US 08/306,473
FILING DATE: 15-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   82.6%; Score 19; DB 6; Length 18; 60.0%; Pred. No. 9.4e+02; tive 0; Mismatches 2; Indels
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STREET: 10 South Wacker Drive, Suite 3000
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                                                                                                                                                                                                                                                                                                                NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0007-999
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELETAX: 650-493-4935
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 199:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
                                                      MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASLSEQ VERSION 2.0
CURRENT APPLICATION DATA:
FILING DATE: 29-OCT-2002
CLASSIFICATION:
                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,136
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
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Best Local Similarity 60.0°
Matches 3; Conservative
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STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTHER INFORMATION: C. US-10-283-599-199
                  ZIP: 10036-2811
COMPUTER READABLE FORM:
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Query Match
Best Local Similarity
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US-08-277-660A-16
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                                                                                  February 12, 2003, 11:12:42; Search time 4.58333 Seconds (without alignments) 32.098 Million cell updates/sec
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Sequence 3, R
Sequence 6, R
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Sequence 29,
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Sequence 1
Sequence 3
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.3
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US-09-435-618-29

US-09-035-618-29

US-08-277-660A-3

US-08-424-957-2

US-08-424-957-2

US-08-424-957-3

US-08-424-957-3

US-09-035-618-2

US-09-035-618-2

US-09-035-618-2

US-09-035-618-1

US-08-424-957-15

US-09-035-618-15

US-08-424-957-15

US-08-428-11

US-08-438-114-15

US-08-709-173-11

US-08-709-173-11

US-08-638-911A-15
                                                                                                                                                                                                                                                                          Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                             262574 segs, 29422922 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                Maximum Match 100%
Listing first 45 summaries
                                                          OM protein - protein search, using sw model
                                                                                                                                                                                                                 Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                           Issued Patents AA:*
                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 08
                                                                                                                                        US-09-403-440A-1
23
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Match Length DB
                                                                                                                                                                                                                                                                                                      Minimum DB seq length: 0
Maximum DB seq length: 25
                                                                                                                                                                         1 PXXLW 5
                                                                                                                                                                                                      BLOSUM62
                                                                                                                                                                                                      Scoring table:
                                                                                                                                                            Perfect score:
                                                                                                                                                                            Sequence:
                                                                                                                                                                                                                                                Searched:
                                                                                                                                                                                                                                                                                                                                                                                                             Database
                                                                                        Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Result
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APPLICANT: Pickaley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
CORRESPONDENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCES:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER: BADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: BM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
CLASSIFICATION:
                                                                                                                                                            9, <u>,</u>
10,
                                                                                                                                                                                   11,
12,
13,
                                                                                                                                 12,
18,
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Sequence 1
Sequence 1
                                                                Sequence
Sequence
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Sequence
Sequence
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                                     Sequence
                         Sequence
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         US-08-709-177-8

US-08-709-177-11

US-08-277-660A-8

US-08-277-660A-8

US-08-277-660A-8

US-08-277-660A-8

US-08-424-957-11

US-08-424-957-11

US-08-035-686-11

US-09-035-686-12

US-09-035-686-12

US-09-035-680-18

US-08-277-660A-10

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11

US-08-277-660A-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACTORNEY/AGENT INPORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: 24,190
REPERENCE/DOCKET NUMBER: 4.60244/WHD
TELECOMMUNICATION INFORMATION:
TELERIONE: (415) 781-189
TELERY: (415) 398-3249
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                    ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                      ; Sequence 16, Application US/08277660A
; Patent No. 5702908
; GENERAL INFORMATION:
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US-08-277-660A-16
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Gaps

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91.3%; Score 21; DB 1; Length 11; 60.0%; Pred. No. 1.5e+02; ative 0; Mismatches 2; Indels

Conservative

linear

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                                                                        Sequence 29, Application US/08424957
Patent No. 5770377
Betent No. 5770377

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STREET: Four Embarcadero Center, Suite 3400
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CITY: San Francisco
STREET: Four Embarcadero Center, Suite 
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Patent No. 615391

APPLICANT: Pickeley, Steven M.
APPLICANT: Lane, David P.

TILLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert

STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: California

COUNTRY: United States
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION: DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGRY INCRMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A-61228/WHD
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TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 791-1999
TELERAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
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94111-4187
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RESULT 2
US-08-424-957-29
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Gaps
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Patent No. 5702908
GENERAL INFORMATION:
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
91.3%; Score 21; DB 4; Length 11;
Best Local Similarity 60.0%; Pred. No. 1.5e+02;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SCFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-UUL-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H:
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET UNMBER: 24,190
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                               NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMNUNICATION INFORMATION:
TELEPAX: (415) 781-1989
TELEFAX: (415) 398-3249
TELEFX: 910 277299
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-ARR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (415) 781-1989
TELEFAX: (415) 388-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy dif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: unknown
US-09-035-686-29
                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 FXXLW 5
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1 FXXLW 5
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Sequence 3, Application US/08277660A

BENERAL INFORMATION:
APPLICANT: Dickeley, Steven M.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Plehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
CITY: Content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the content of the con
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
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87.0%; Score 20; DB 1; Length 6;
Best Local Similarity 60.0%; Pred. No. 2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: COMPUTER: COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILLING DALE: ACCOUNTED TO THE STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF STATEMENT OF 
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US-08-424-957-2
; Sequence 2, Application US/08424957
; Patent No. 57/0377
; GENERAL INPORMATION:
; APPLICANT: Picksley, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
SEQUENCE CHARACTERISTICS
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                                                                                         LENGTH: 6 amino acids
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                                                                                                                                   TYPE: amino acid STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                 linear
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US-08-277-660A-2
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US-08-277-660A-3
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PRICAMIN Lano, David P.

TITLE OF INVERTION: Interruption of Binding of WDMA and P53
TITLE OF INVERTION: Interruption of Binding of WDMA and P53
TITLE OF INVERTION: Protein and Therapeutic Application Thereof
NUMBERS OF SEQUENCES: 50
CONTRET. David States
SOUTH TITLE OF INVERTION: Protein and Therapeutic Application Thereof
COUNTY: United States
COUNTY: United States
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APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF EXQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 87.0%; Score 20; DB 1; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
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ZIP: 94111-4187
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPAtible
COMPUTER: IBM PC COMPAtible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATION
SOFTWARE: PATENTIAN DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTONEY/AGRAT INFORMATION:
MANNEY/AGRAT INFORMATION:
                   PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: DEGGEV, Walter H.
REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: A-61228/WHD
TELECOWMUNICATION INFORMATION:
TELEPHONE: (415) 398-3249
TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
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REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-424-957-6; Sequence 6, Application US/08424957; Patent No. 5770377
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TELEFAX: (415) 38-3249
TELEX: 910 277299
INPORMATION FOR SEQ 1D NO: 6: SEQUENCE CHARACTERISTICS: ENGTH: 6 amino acids
TYPE: amino acid STRANDEDNESS:
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US-08-424-957-3
                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS:
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CLASSIFICATION:
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APPLICANT: Picksley, Steven M.
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
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Query Match 87.0%; Score 20; DB 1; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
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APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY, AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERRACE/DOCKET NUMBER: 3-61228/WHD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                            ; Sequence 2, Application US/09035686
; Patent No. 6153391
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US-03-035-686-3
Sequence 3, Application US/09035686
Patent No. 6153391
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: Four Embarcadero
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
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Matches 3; Conservative
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TYPE: amino acid
STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: unknown
US-09-035-686-2
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                                                                                             1 FXXLW 5
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US-09-035-686-2
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TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
STORMER: PATENTIN Release #1.0, Version #1.30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87.0%; Score 20; DB 4; Length 6; 60.0%; Pred. No. 2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781,1989
                                                                                                                                                                                                                                                                                                                CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION UNDERE: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 6 amino acids
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Best Local Similarity 60.0
Matches 3; Conservative
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US-09-035-686-3
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APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
TITLE OF INVENTION: ANTIBODIES
CORRESPONDENCES: 27
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; Indels
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MEDIUM TYPE: Diskette
COMPUTER: IBM COMPALIALE
COMPUTER: IBM COMPALIALE
OFFRATING SYSTEM: Windows
SOFTWARE: FASTENG for Windows
SOFTWARE: FASTENG for Windows
SOFTWARE: L2-MAY-1998
CLASSIFICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 87.0%; Score 20; DB 4; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches
            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-ARR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH : 6 anino acids
                                                                                                                                                                                                                             A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 12
US-09-081-975-1
Sequence 1, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 02110
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TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
                                                                           STREET: Four Embarcacco
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              United States
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 amino acids
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                       NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FXXLW 5
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US-08-424-957-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Interruption of Binding of MDM2 and P53 TITLE OF INVENTION: Protein and Therapeutic Application Thereof ORRESPONDENCEs. 27 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
0
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Patent No. 5770377
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
                                                                                              Query Match 87.0%; Score 20; DB 4; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       87.0%; Score 20; DB 1; Length 7; 60.0%; Pred. No. 2e+05; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMONIALCATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                   Sequence 27, Application US/08277660A
Patent No. 5702908
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             California
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Best Local Similarity
'... 3; Conserve
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US-08-277-660A-27
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US-08-424-957-15
                                                        US-09-081-975-1
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Gaps
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APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .
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                                                                            ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STRET: Pour Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
FILLING DATE: US/08/424,957
FILLING DATE: 19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE: US/08/424,957
FILING DATE: 19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 19-APR-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-ULL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 19, Application US/08424957; Patent No. 5770377; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFRENCE/DOCKET NUMBER: A-61;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
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### APPLICATION NUMBER: US 08/277,660
### FILING DATE: 20-JUL-1994
### ATTOREX FACENT INFORMATION:
### NAME: Dreger, Walter H.
### REGISTRATION NUMBER: 24,190
### REGISTRATION NUMBER: 24,190
### REFRENCENCY NUMBER: A-61228/WHD
### TELEPHONE: (415) 78-1989
### TELEPHONE: (415) 78-1989
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### TELEPHONE: (415)
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